

Carbohydrat Metabolism



گلوکز

پروات ← لاکتات

استیل کوآنزیم A

اجسام کتونی

کلسترول

چرخه کربس

اسید چرب

تری گلیسرید

- گلیکولیز
- گلیکوژنز
- گلیکوژنولیز
- مسیر پنتوز فسفات
- گلوکونئوژنز
- لیپوژنز
- لیپولیز



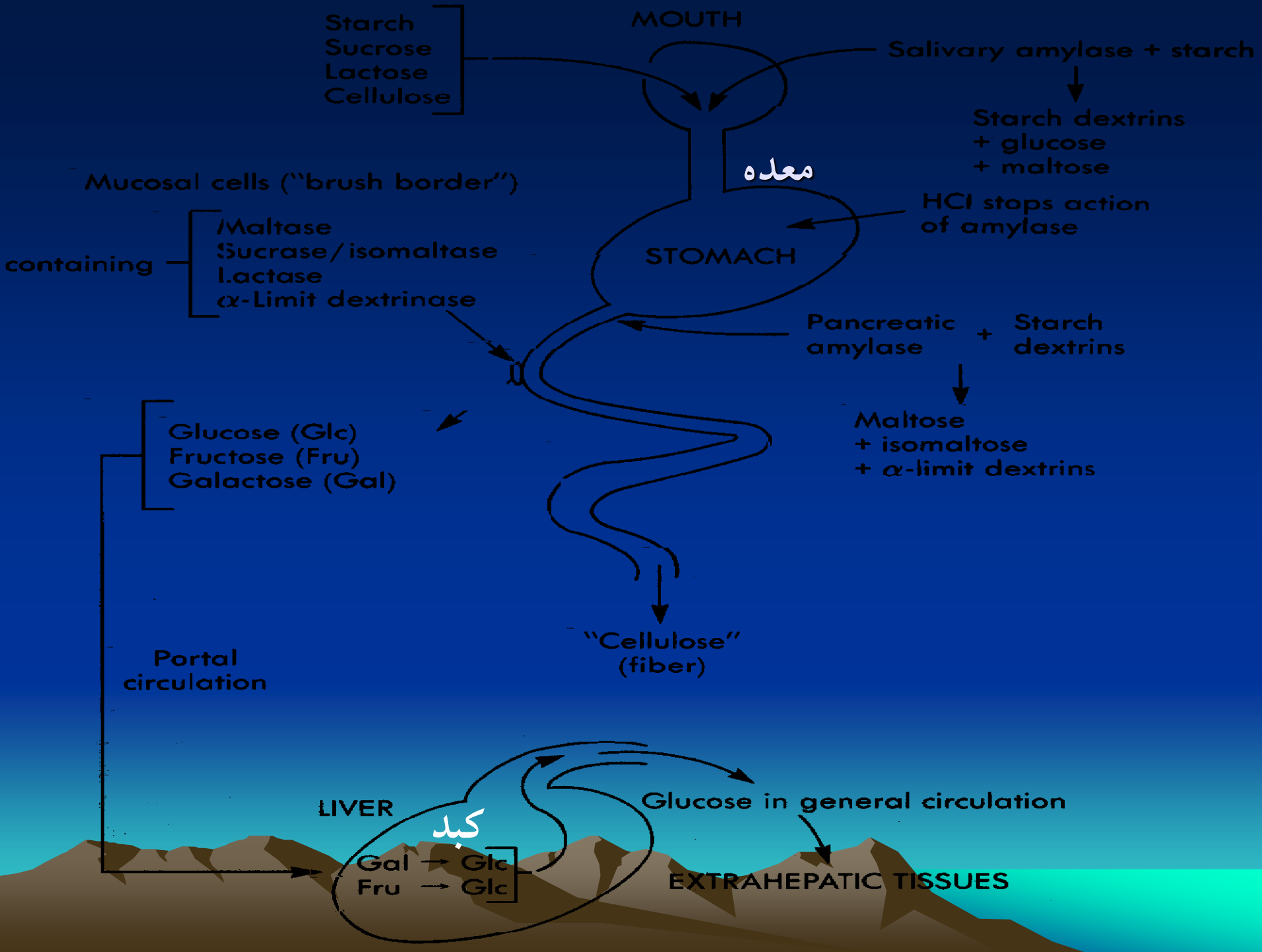
Digestion & Carbohydrate Absorption



Table 19–2. Glucose transporters.

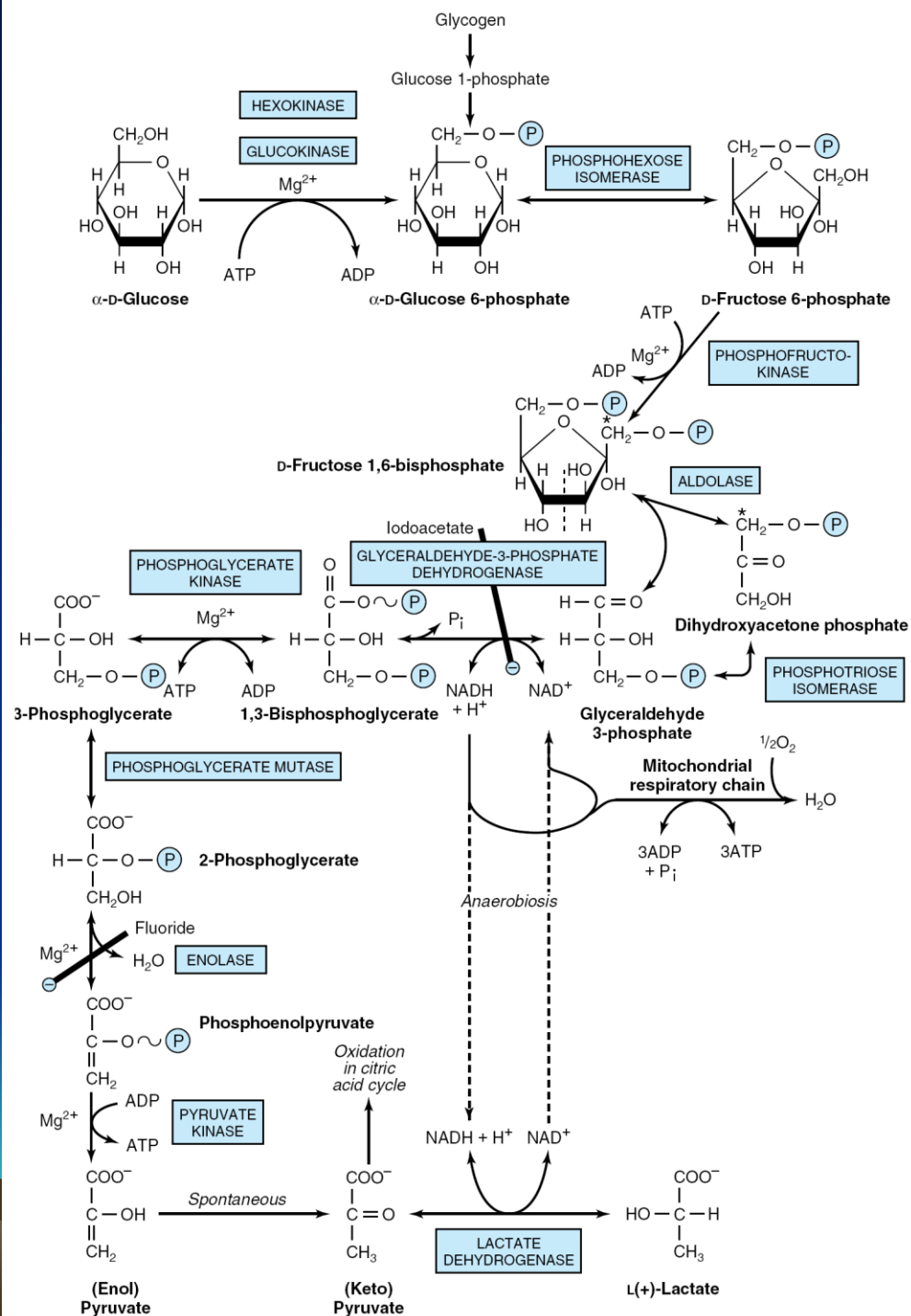
	Tissue Location	Functions
Facilitative bidirectional transporters		
GLUT 1	Brain, kidney, colon, placenta, erythrocyte	Uptake of glucose
GLUT 2	Liver, pancreatic B cell, small intestine, kidney	Rapid uptake and release of glucose
GLUT 3	Brain, kidney, placenta	Uptake of glucose
GLUT 4	Heart and skeletal muscle, adipose tissue	Insulin-stimulated uptake of glucose
GLUT 5	Small intestine	Absorption of glucose
Sodium-dependent unidirectional transporter		
SGLT 1	Small intestine and kidney	Active uptake of glucose from lumen of intestine and reabsorption of glucose in proximal tubule of kidney against a concentration gradient



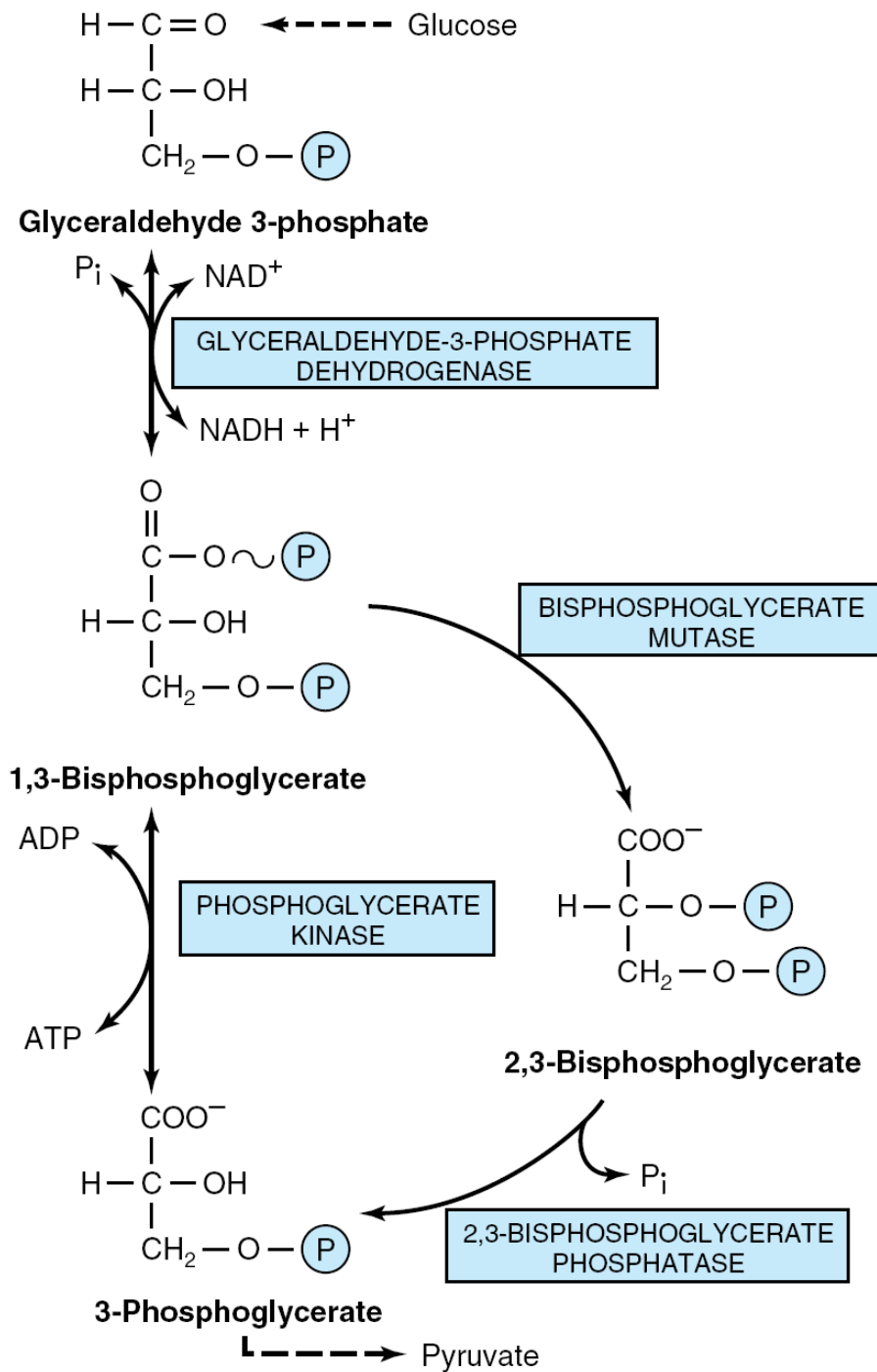


گلیکولیز



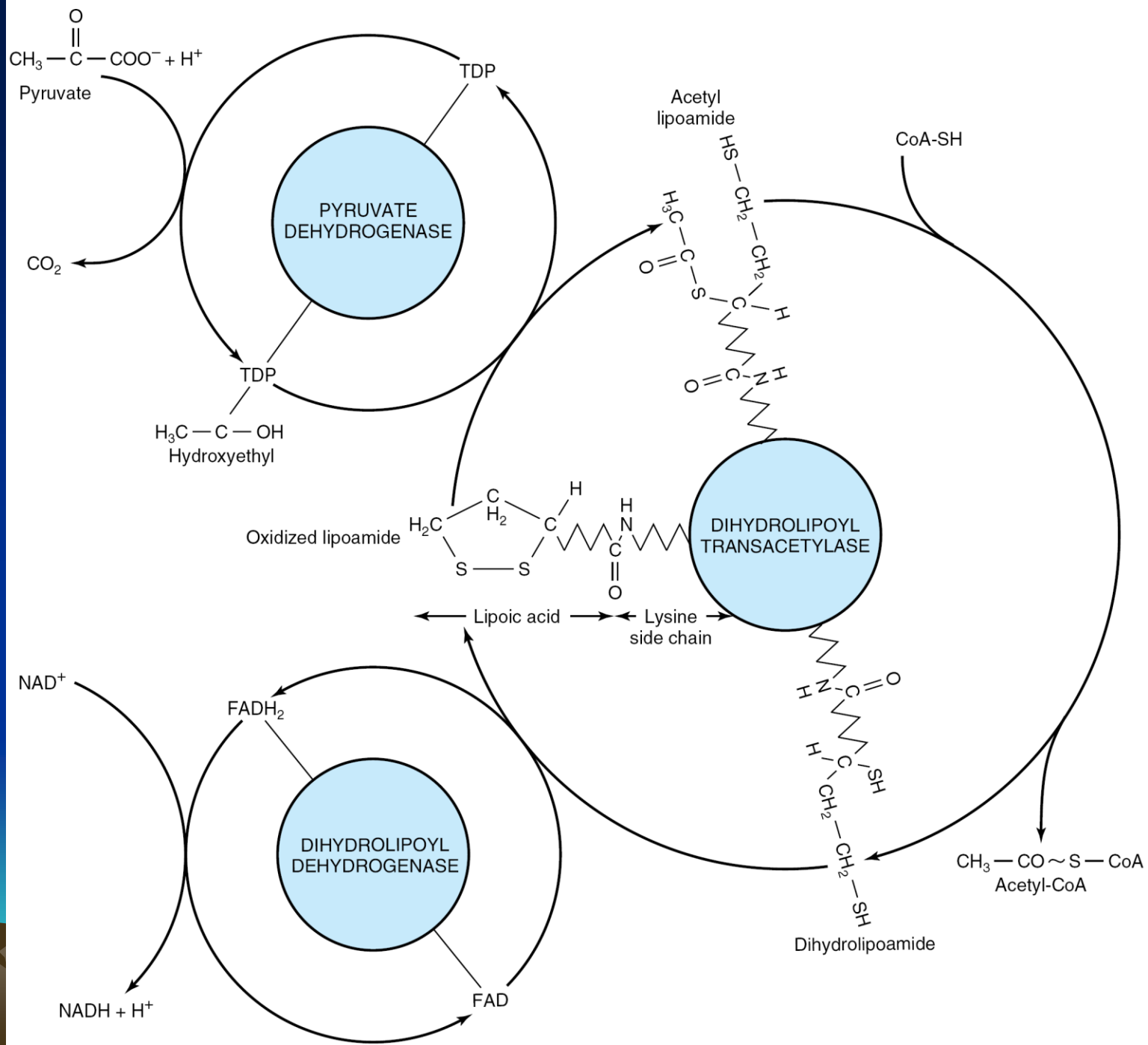


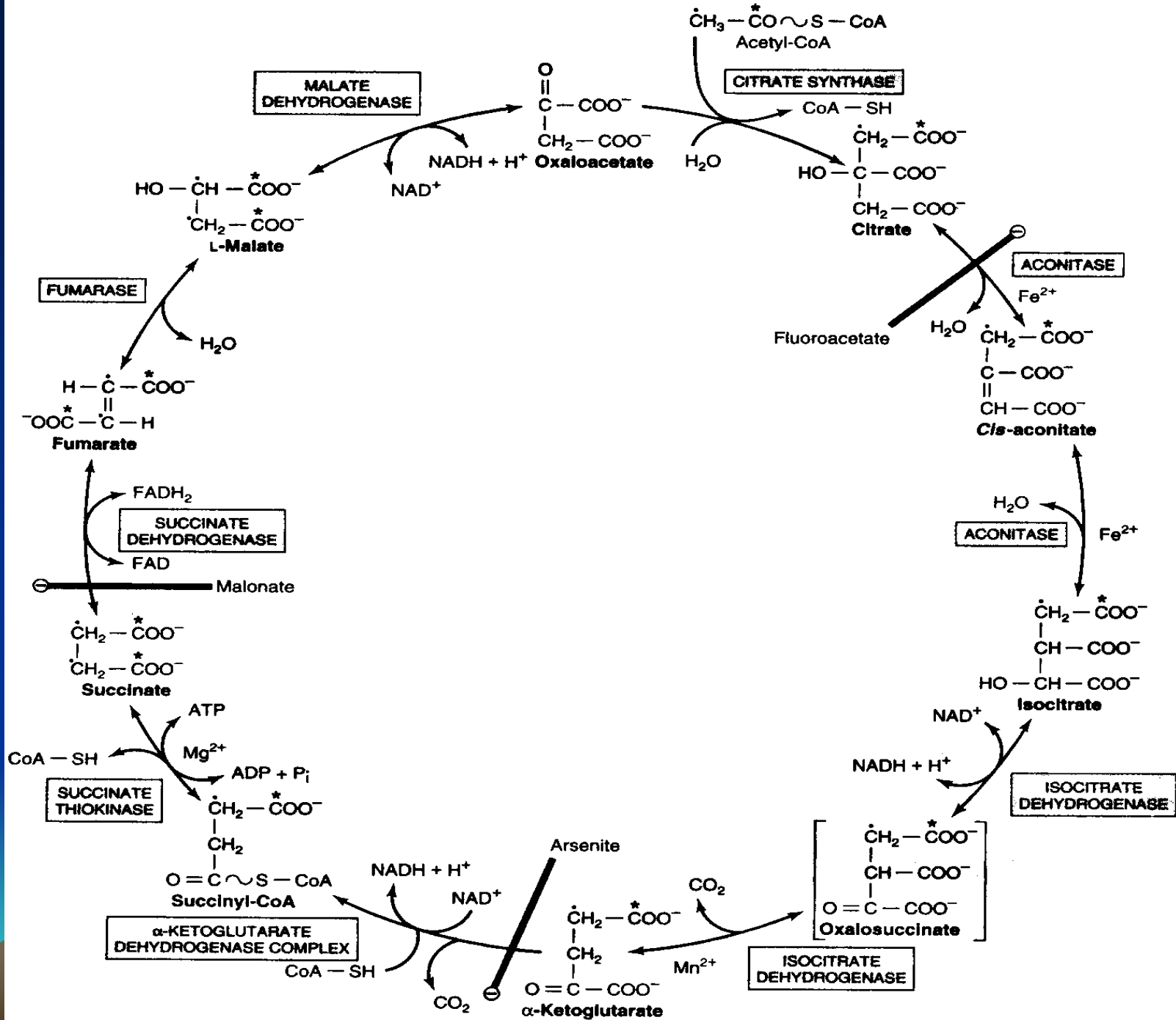
Rapaport Shunt

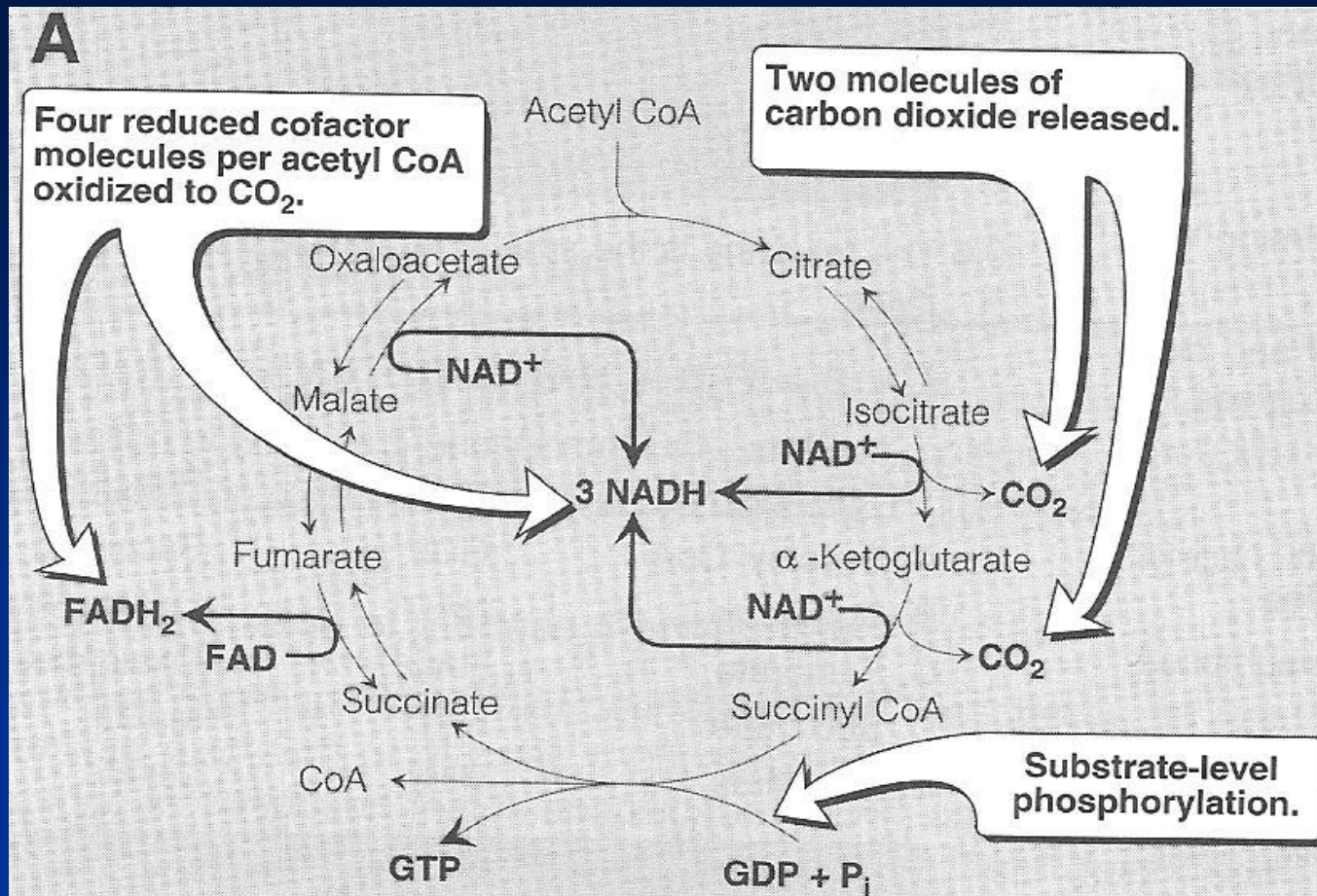


تبدیل پیرووات به استیل کوآنزیم A

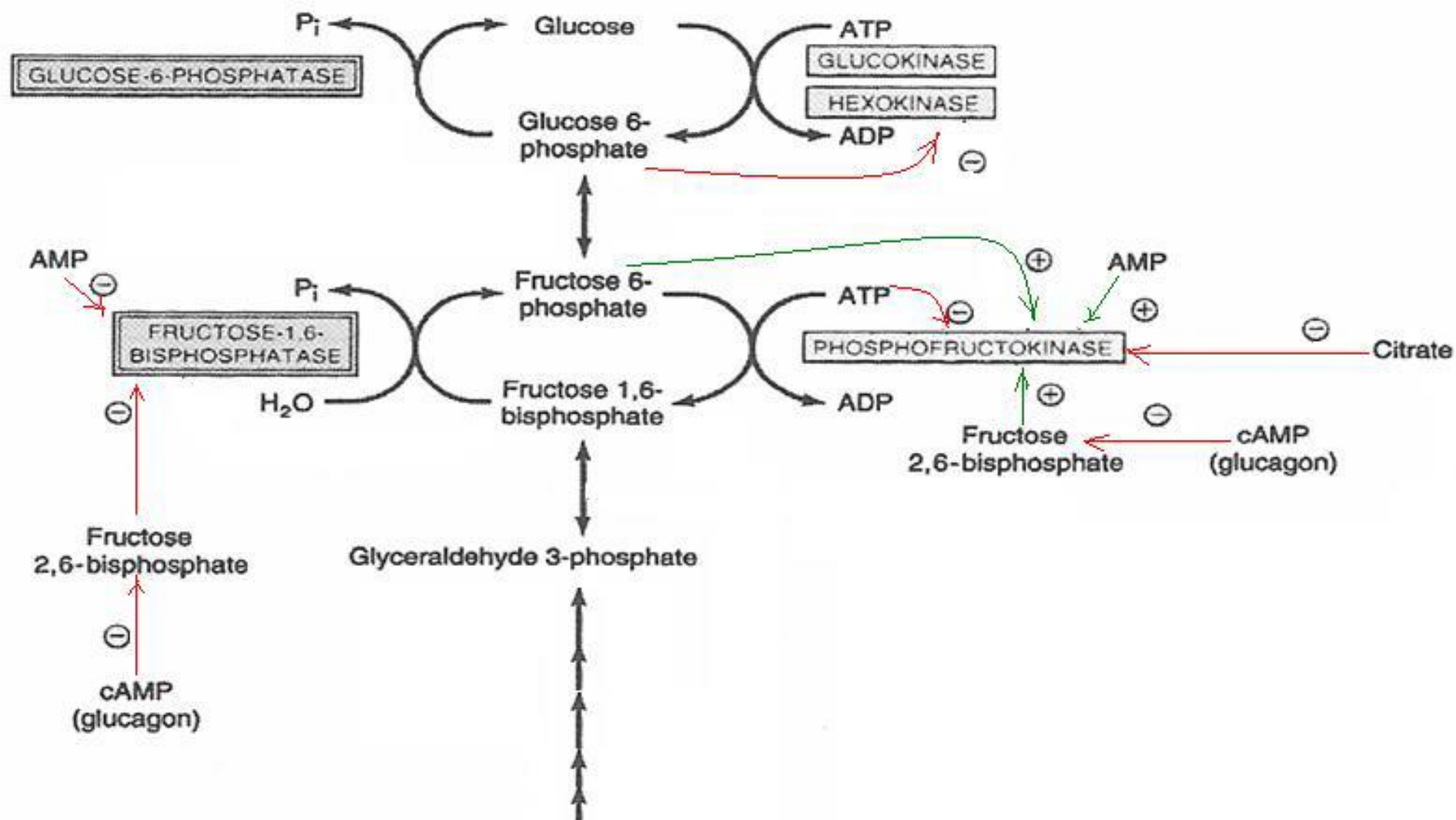


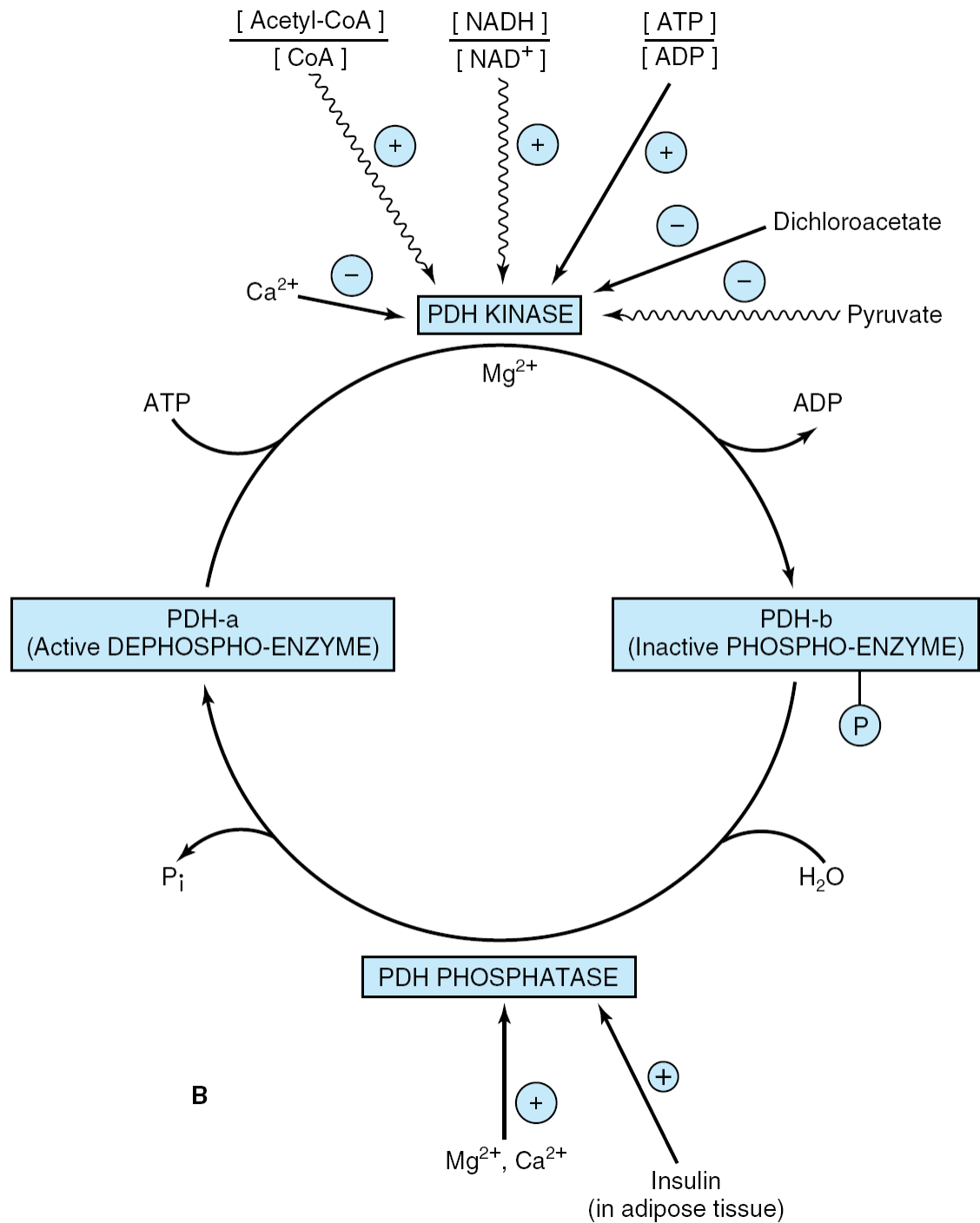
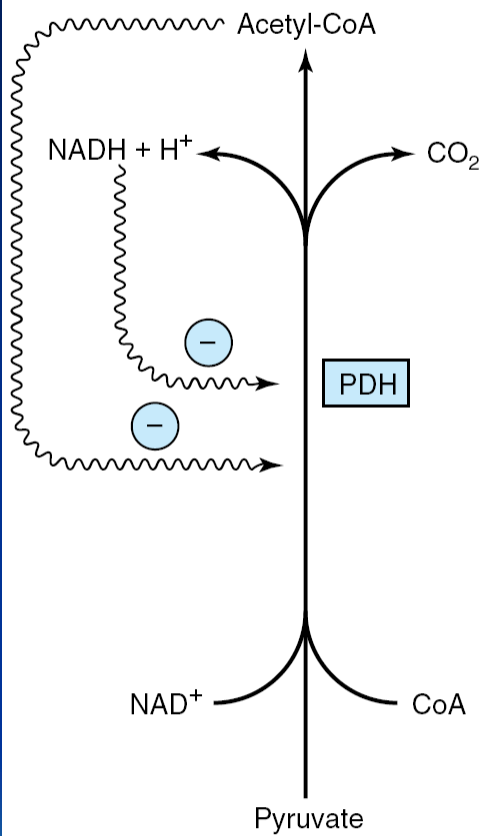


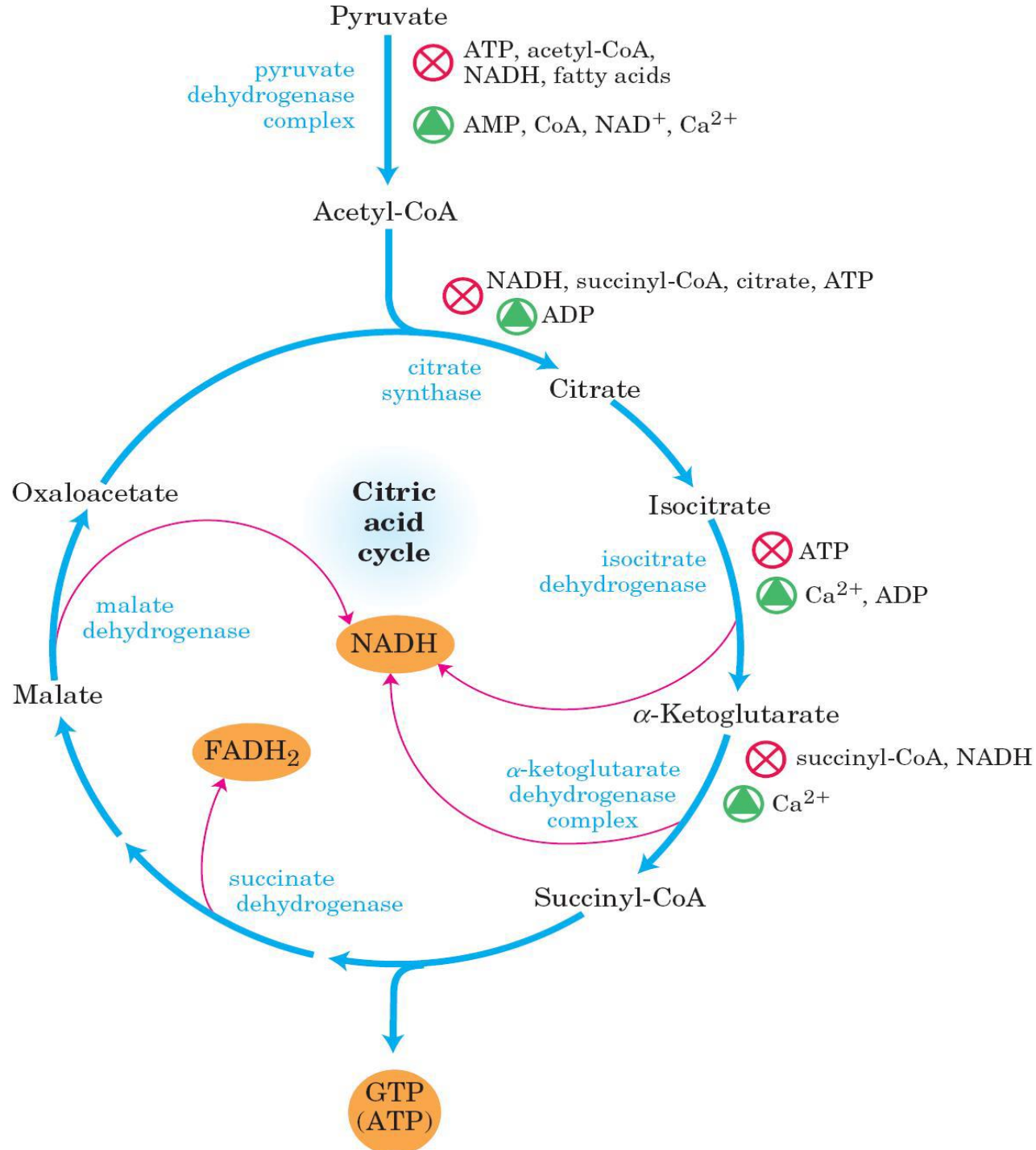




Energy-producing reaction	Number of ATP produced
$3 \text{ NADH} \longrightarrow 3 \text{ NAD}^+$	9
$\text{FADH}_2 \longrightarrow \text{FAD}$	2
$\text{GDP} + \text{P}_i \longrightarrow \text{GTP}$	1
<hr/>	
12 ATP/acetyl CoA oxidized	

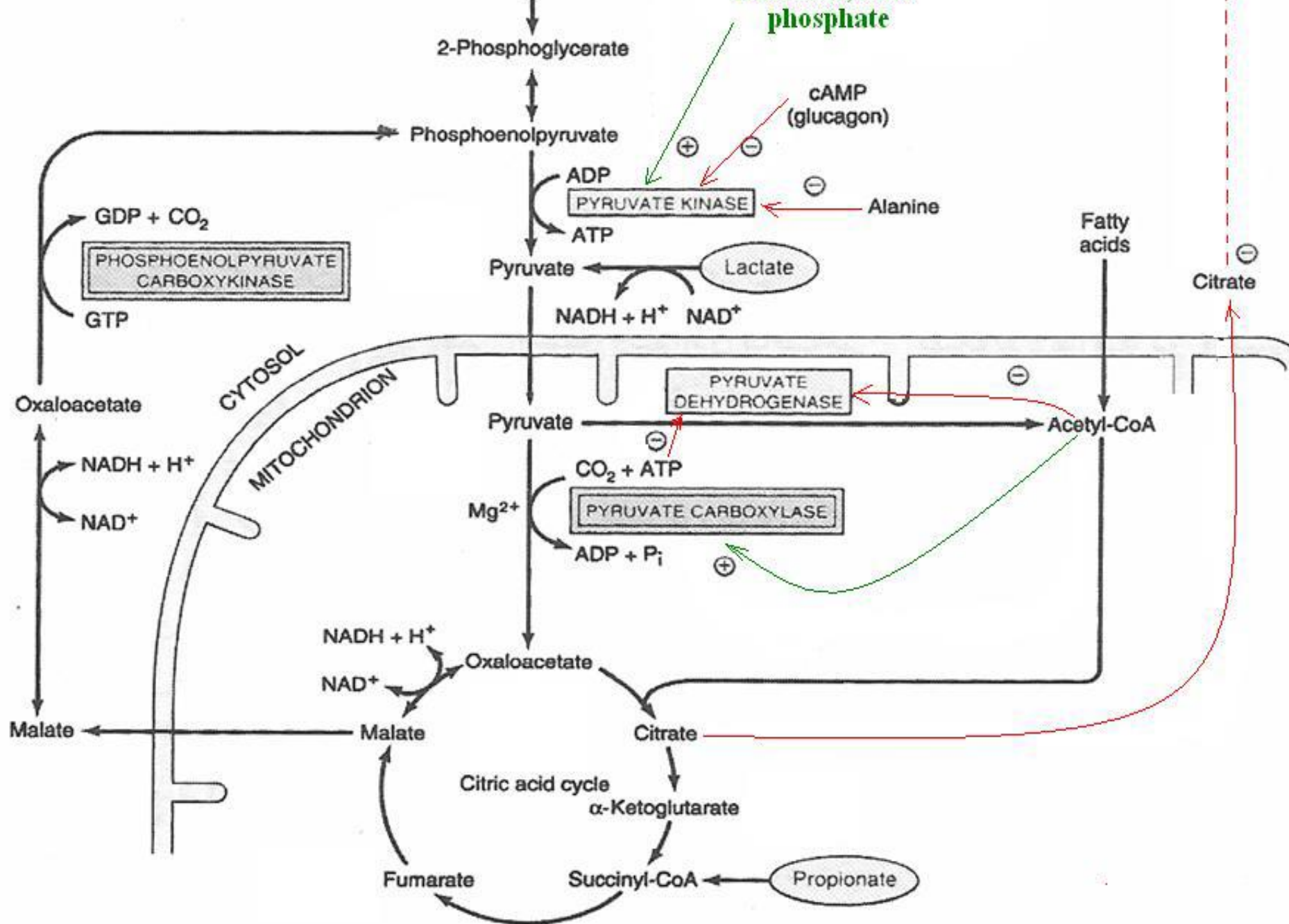


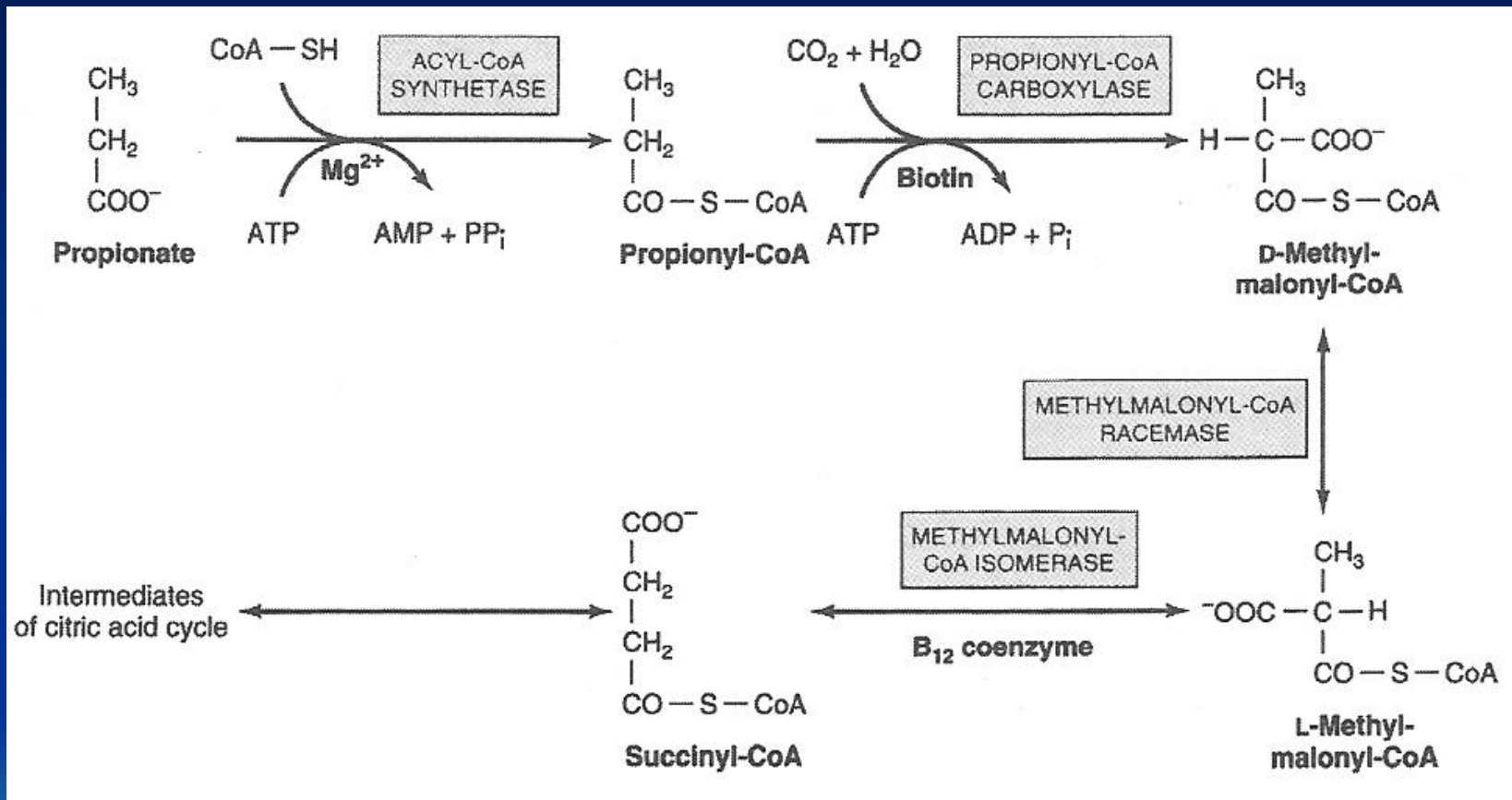


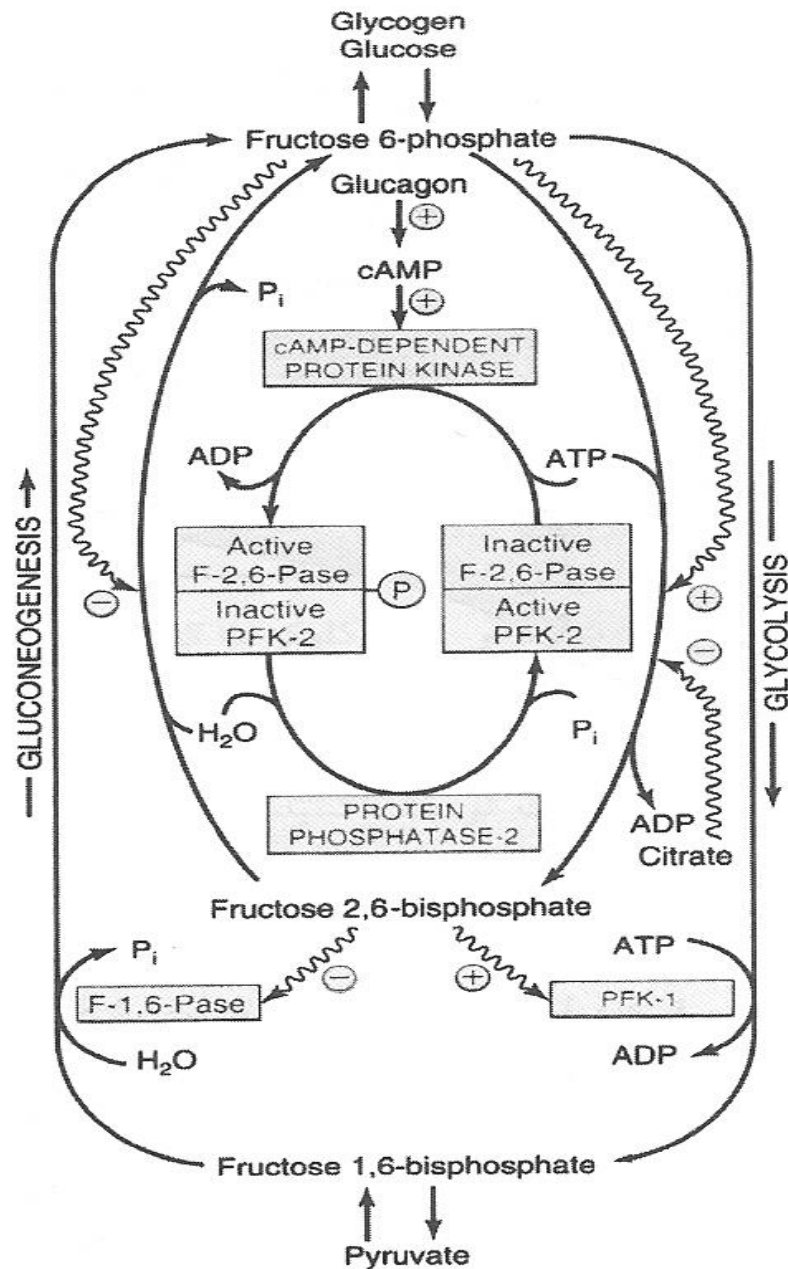


Gluconeogenesis









شکل ۳-۱۹. کنترل گلیکولیز و گلوکونئوژنز در کبد توسط فروکتو

۶،۲-بیس فسفات و آنزیم دوکاره PFK/F-2,6-Pase (۶-فسفوفروکتو-۲-کیناز

۶،۲-کیناز، PFK-1) (۶-فسفاتاز، F-2,6-Pase) و پروتئین فسفاتاز-۲.

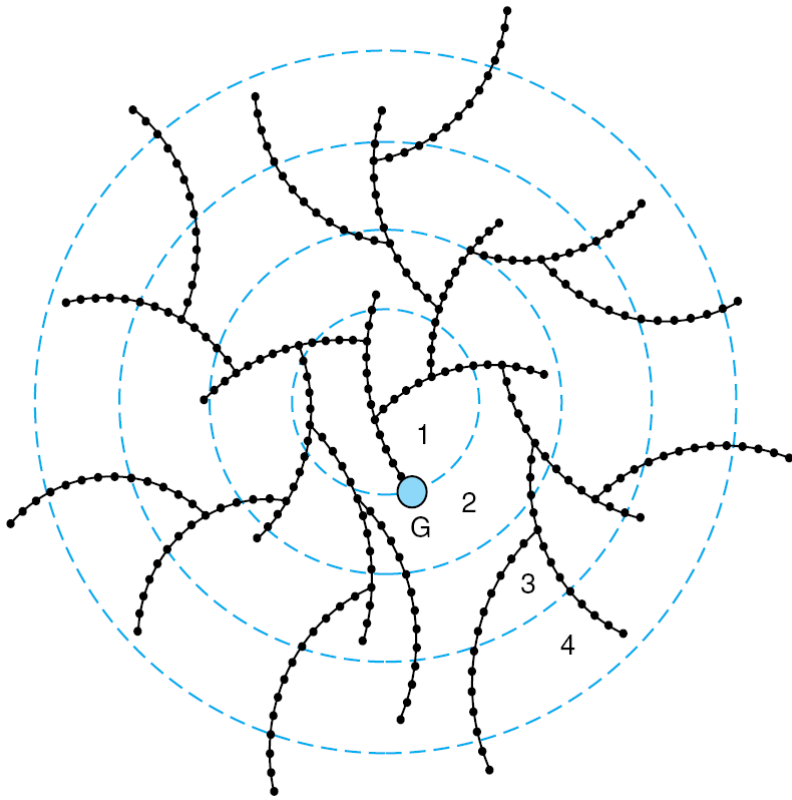
Table 19–1. Regulatory and adaptive enzymes of the rat (mainly liver).

	Activity In		Inducer	Repressor	Activator	Inhibitor
	Carbo- hydrate Feeding	Starva- tion and Diabetes				
Enzymes of glycogenesis, glycolysis, and pyruvate oxidation						
Glycogen synthase system	↑	↓	Insulin		Insulin Glucose 6-phosphate ¹	Glucagon (cAMP) phos- phorylase, glycogen
Hexokinase						Glucose 6-phosphate ¹
Glucokinase	↑	↓	Insulin	Glucagon (cAMP)		
Phosphofructokinase-1	↑	↓	Insulin	Glucagon (cAMP)	AMP, fructose 6-phosphate, P _i , fruc- tose 2,6-bisphos- phate ¹	Citrate (fatty acids, ketone bodies), ¹ ATP, ¹ glucagon (cAMP)
Pyruvate kinase	↑	↓	Insulin, fructose	Glucagon (cAMP)	Fructose 1,6- bisphosphate ¹ , in- sulin	ATP, alanine, glucagon (cAMP), epinephrine
Pyruvate dehydro- genase	↑	↓			CoA, NAD ⁺ , insu- lin, ² ADP, pyruvate	Acetyl-CoA, NADH, ATP (fatty acids, ketone bodies)
Enzymes of gluconeogenesis						
Pyruvate carboxylase	↓	↑	Glucocorticoids, glucagon, epi- nephrine (cAMP)	Insulin	Acetyl-CoA ¹	ADP ¹
Phosphoenolpyruvate carboxykinase	↓	↑	Glucocorticoids, glucagon, epi- nephrine (cAMP)	Insulin	Glucagon?	
Fructose-1,6- bisphosphatase	↓	↑	Glucocorticoids, glucagon, epi- nephrine (cAMP)	Insulin	Glucagon (cAMP)	Fructose 1,6- bisphosphate, AMP, fructose 2,6-bisphos- phate ¹
Glucose-6-phosphatase	↓	↑	Glucocorticoids, glucagon, epi- nephrine (cAMP)	Insulin		

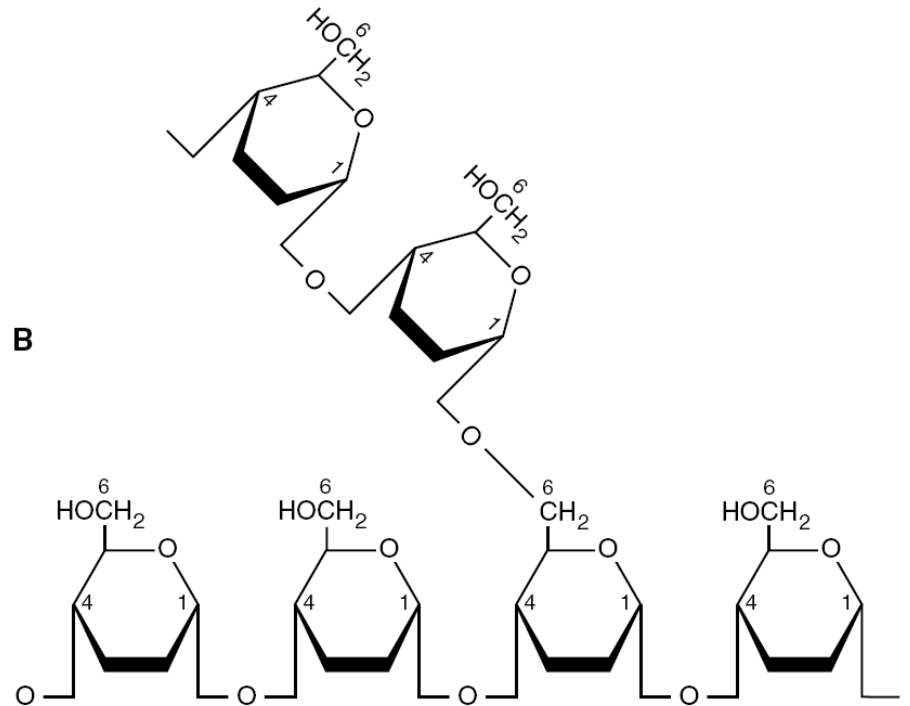
Glycogenesis



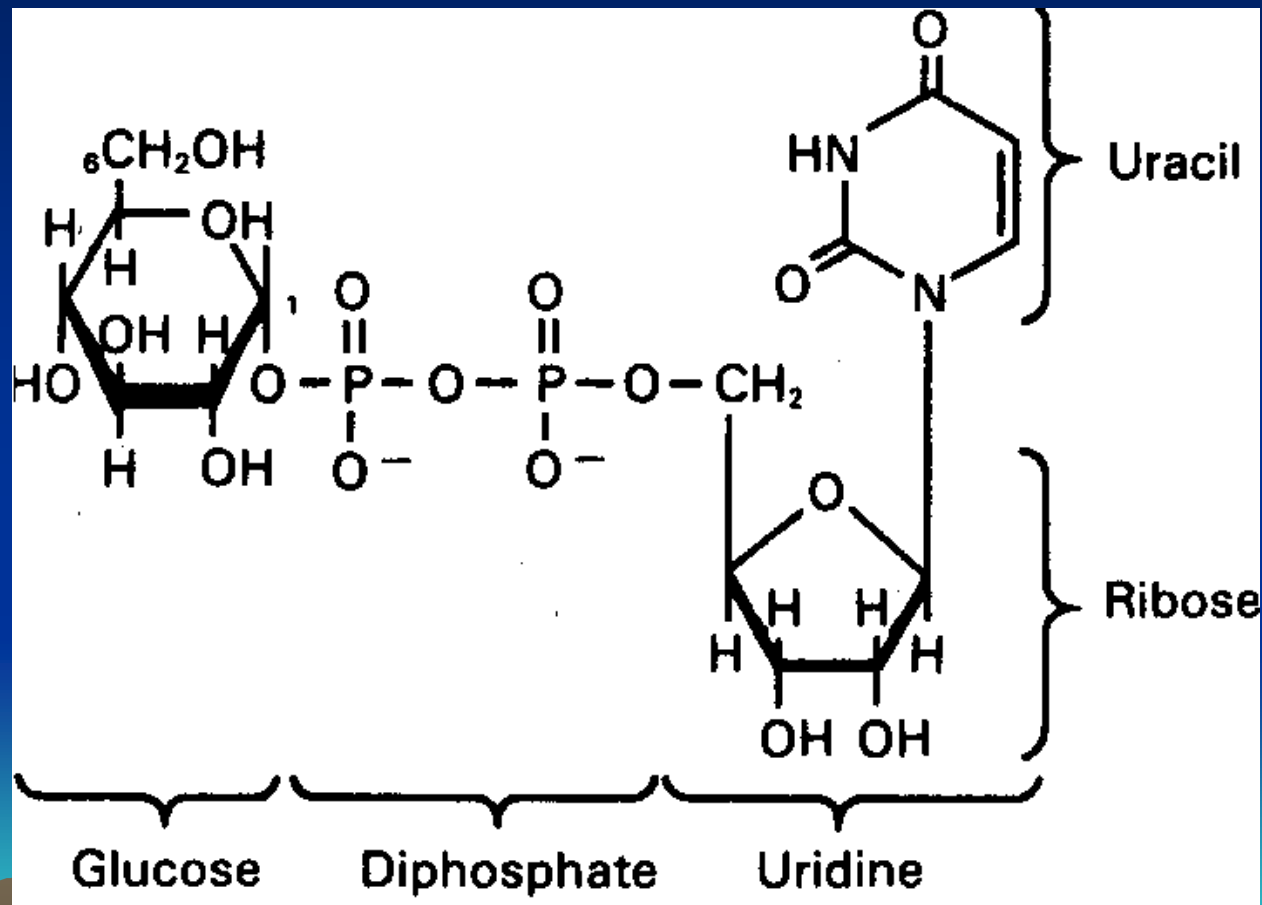
Glycogen molecule

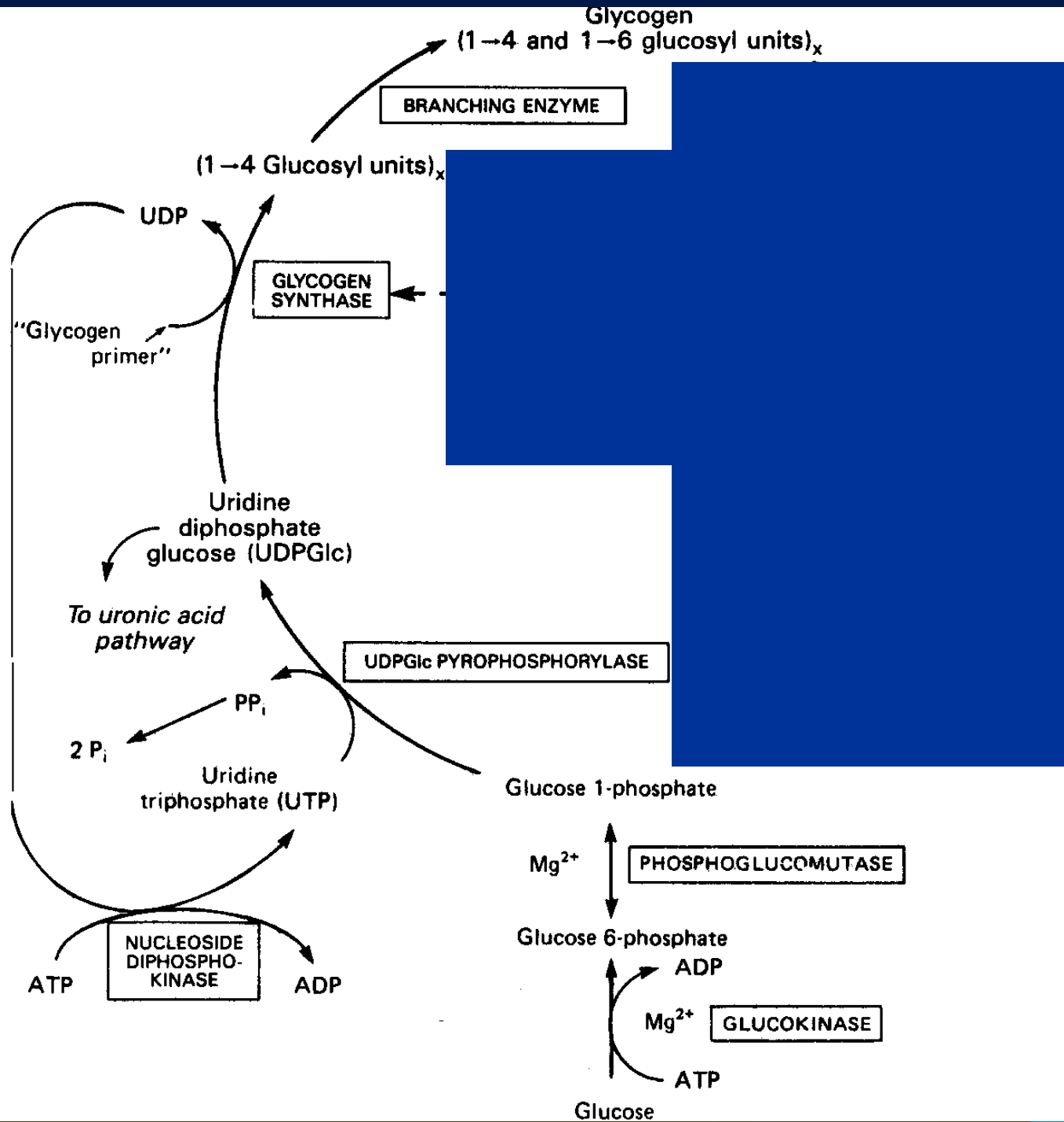


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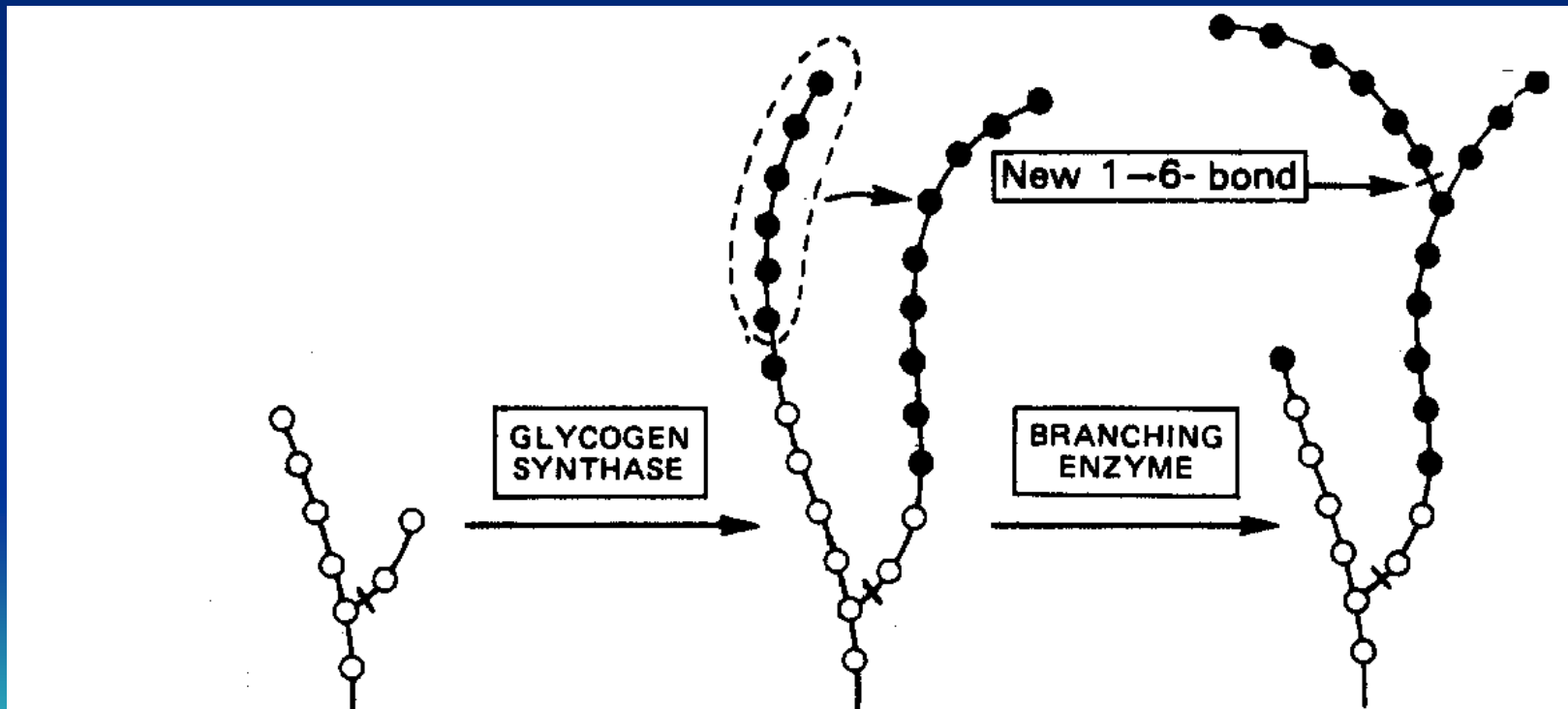


UDP - Glucose





Branching Enzyme



Glycogenolysis



Glycogen
(1→4 and 1→6 glucosyl units)_x

گلیکوژن

P_i

PHOSPHORYLASE

Glucose 1-phosphate

Mg²⁺

PHOSPHOGLUCOMUTASE

Glucose 6-phosphate

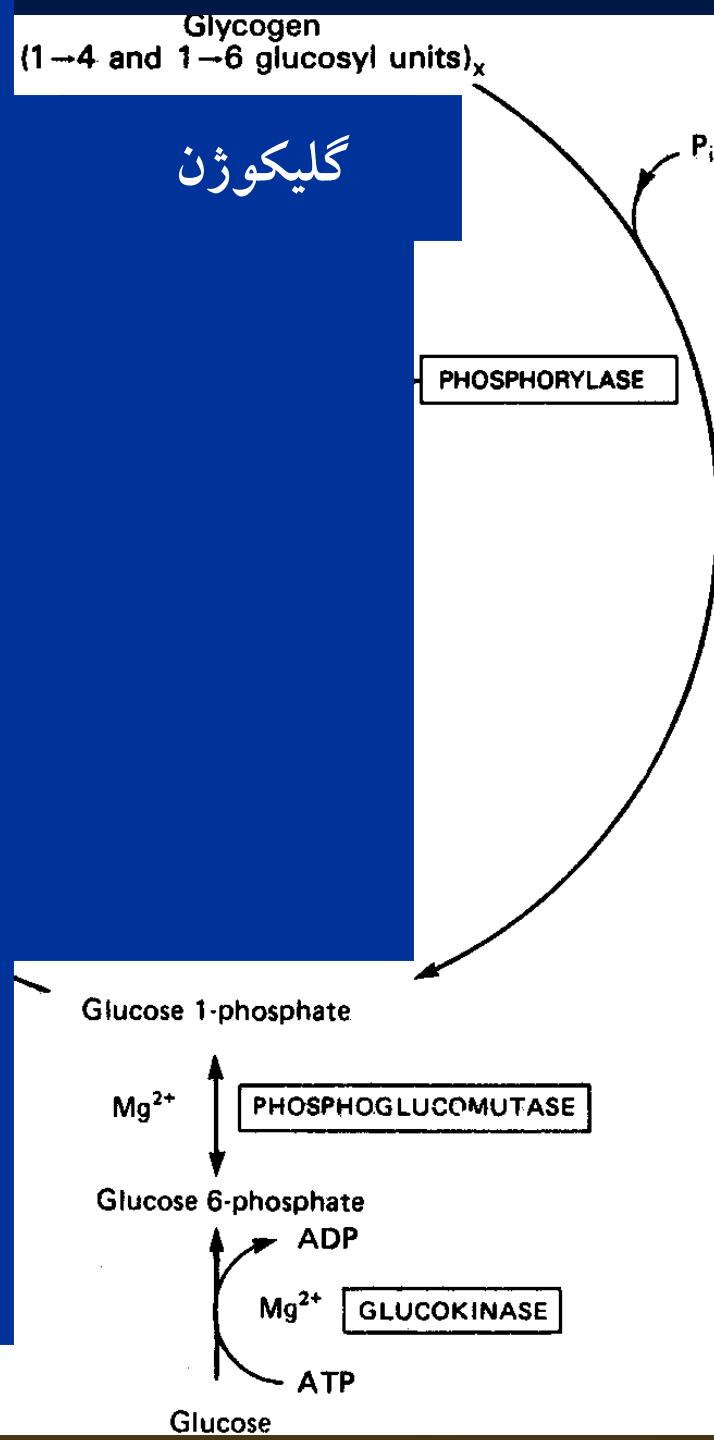
ADP

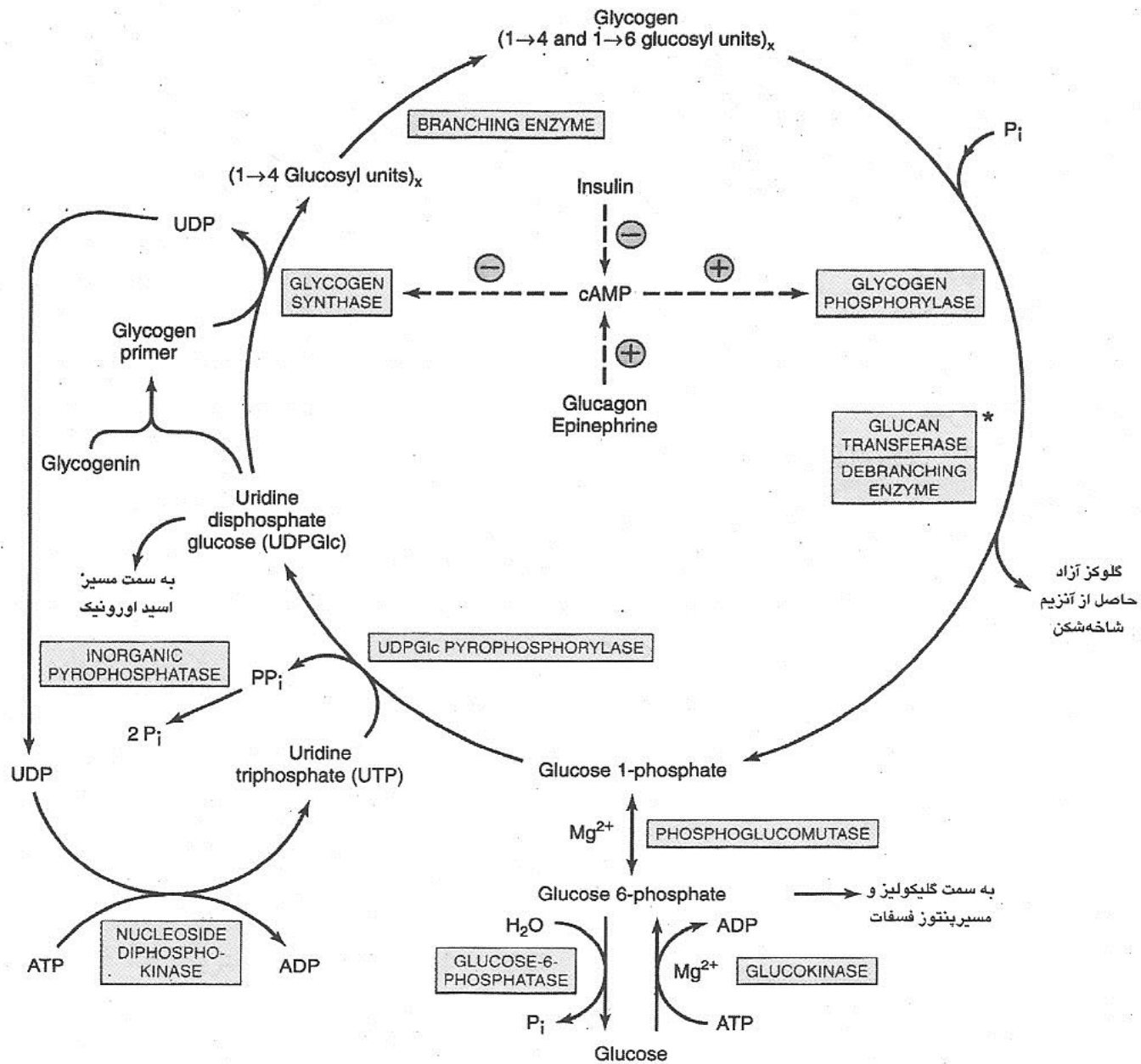
Mg²⁺

GLUCOKINASE

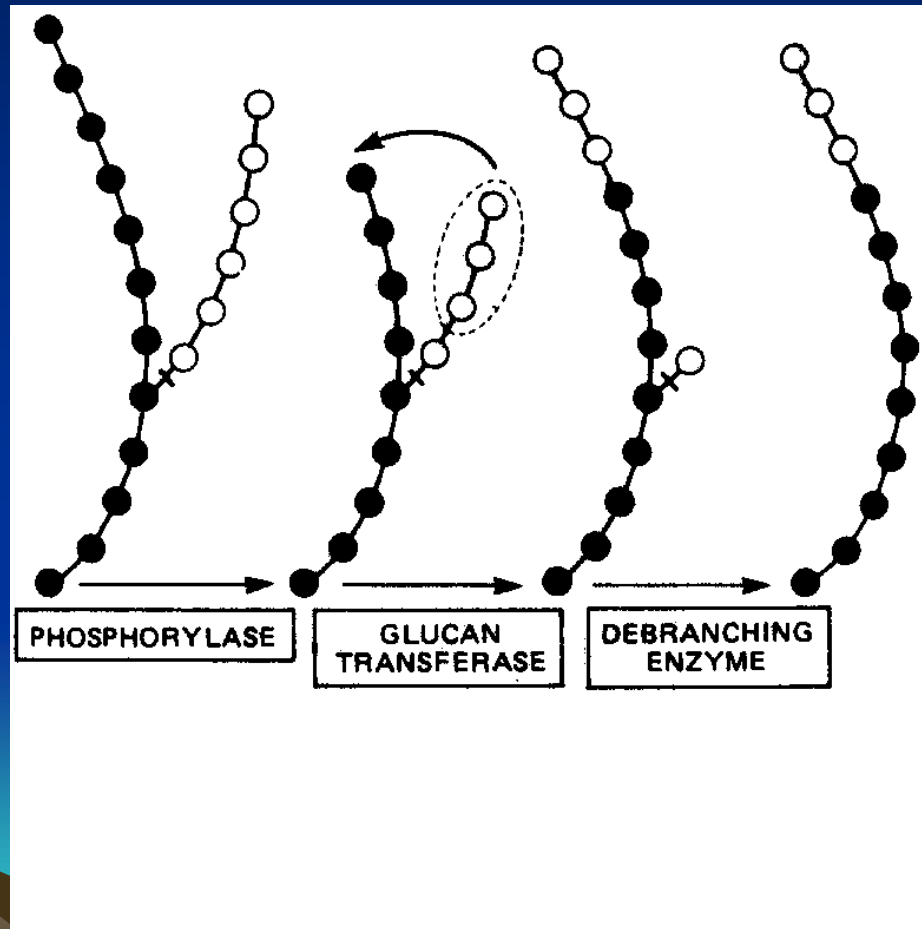
ATP

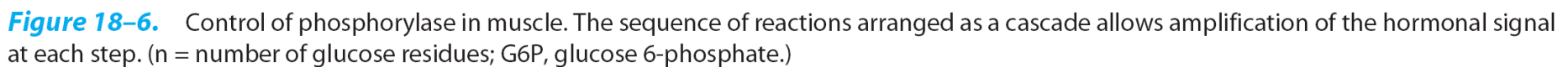
Glucose





Debranching Enzyme





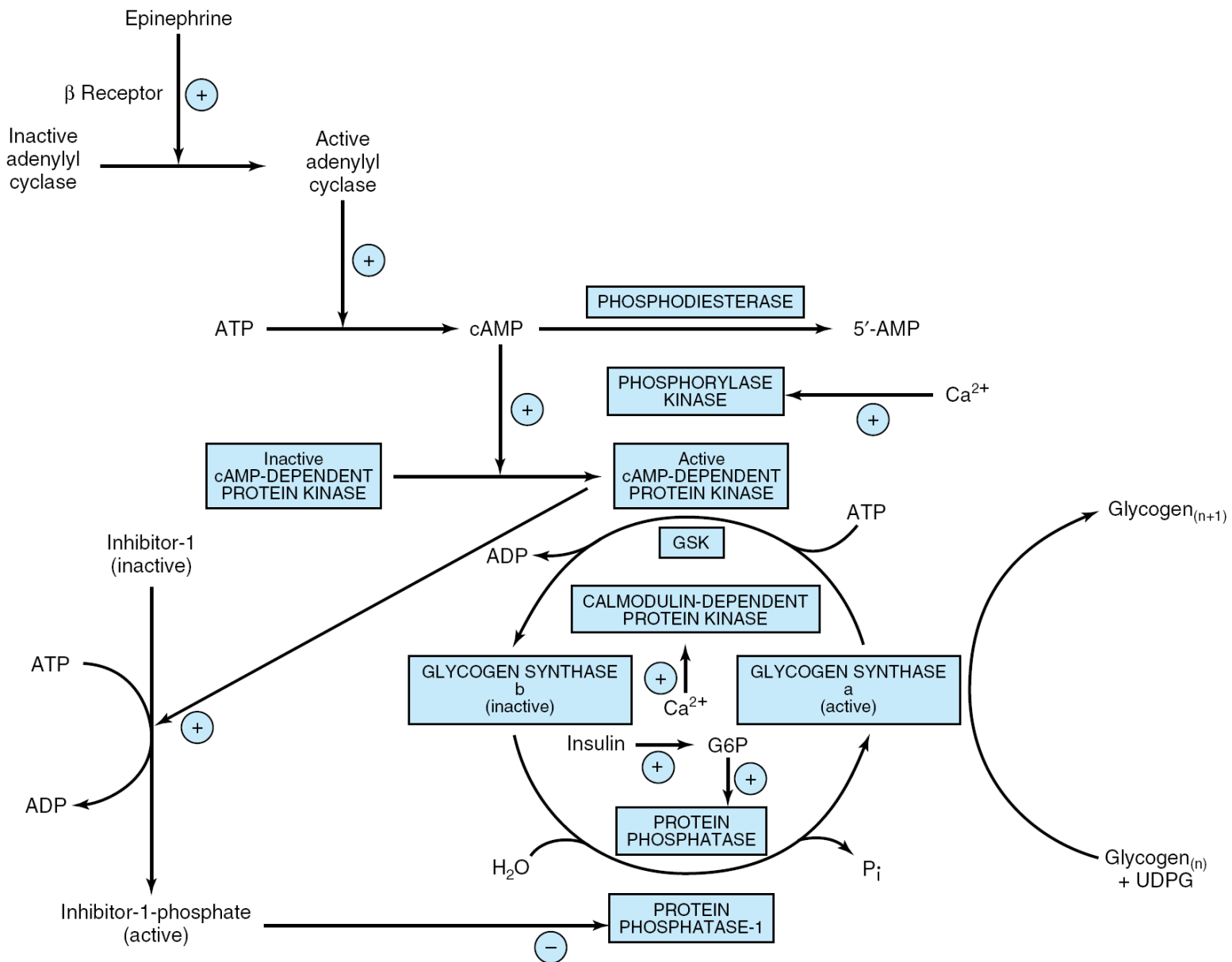


Figure 18–7. Control of glycogen synthase in muscle (n = number of glucose residues). The sequence of reactions arranged in a cascade causes amplification at each step, allowing only nanomole quantities of hormone to cause major changes in glycogen concentration. (GSK, glycogen synthase kinase-3, -4, and -5; G6P, glucose 6-phosphate.)

Glycogenosis

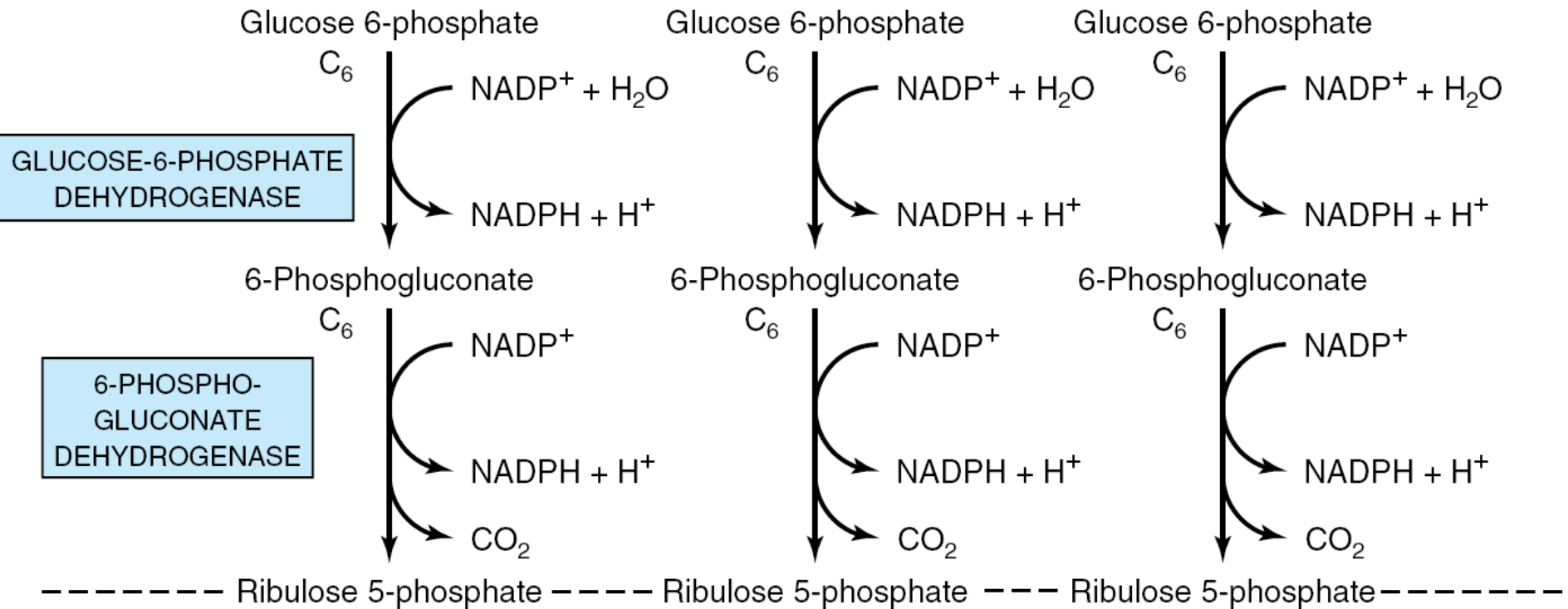
Table 18–2. Glycogen storage diseases.

Glycogenosis	Name	Cause of Disorder	Characteristics
Type I	Von Gierke's disease	Deficiency of glucose-6-phosphatase	Liver cells and renal tubule cells loaded with glycogen. Hypoglycemia, lactic-acidemia, ketosis, hyperlipemia.
Type II	Pompe's disease	Deficiency of lysosomal α -1 \rightarrow 4- and 1 \rightarrow 6-glucosidase (acid maltase)	Fatal, accumulation of glycogen in lysosomes, heart failure.
Type III	Limit dextrinosis, Forbes' or Cori's disease	Absence of debranching enzyme	Accumulation of a characteristic branched polysaccharide.
Type IV	Amylopectinosis, Andersen's disease	Absence of branching enzyme	Accumulation of a polysaccharide having few branch points. Death due to cardiac or liver failure in first year of life.
Type V	Myophosphorylase deficiency, McArdle's syndrome	Absence of muscle phosphorylase	Diminished exercise tolerance; muscles have abnormally high glycogen content (2.5–4.1%). Little or no lactate in blood after exercise.
Type VI	Hers' disease	Deficiency of liver phosphorylase	High glycogen content in liver, tendency toward hypoglycemia.
Type VII	Tarui's disease	Deficiency of phosphofructokinase in muscle and erythrocytes	As for type V but also possibility of hemolytic anemia.
Type VIII		Deficiency of liver phosphorylase kinase	As for type VI.

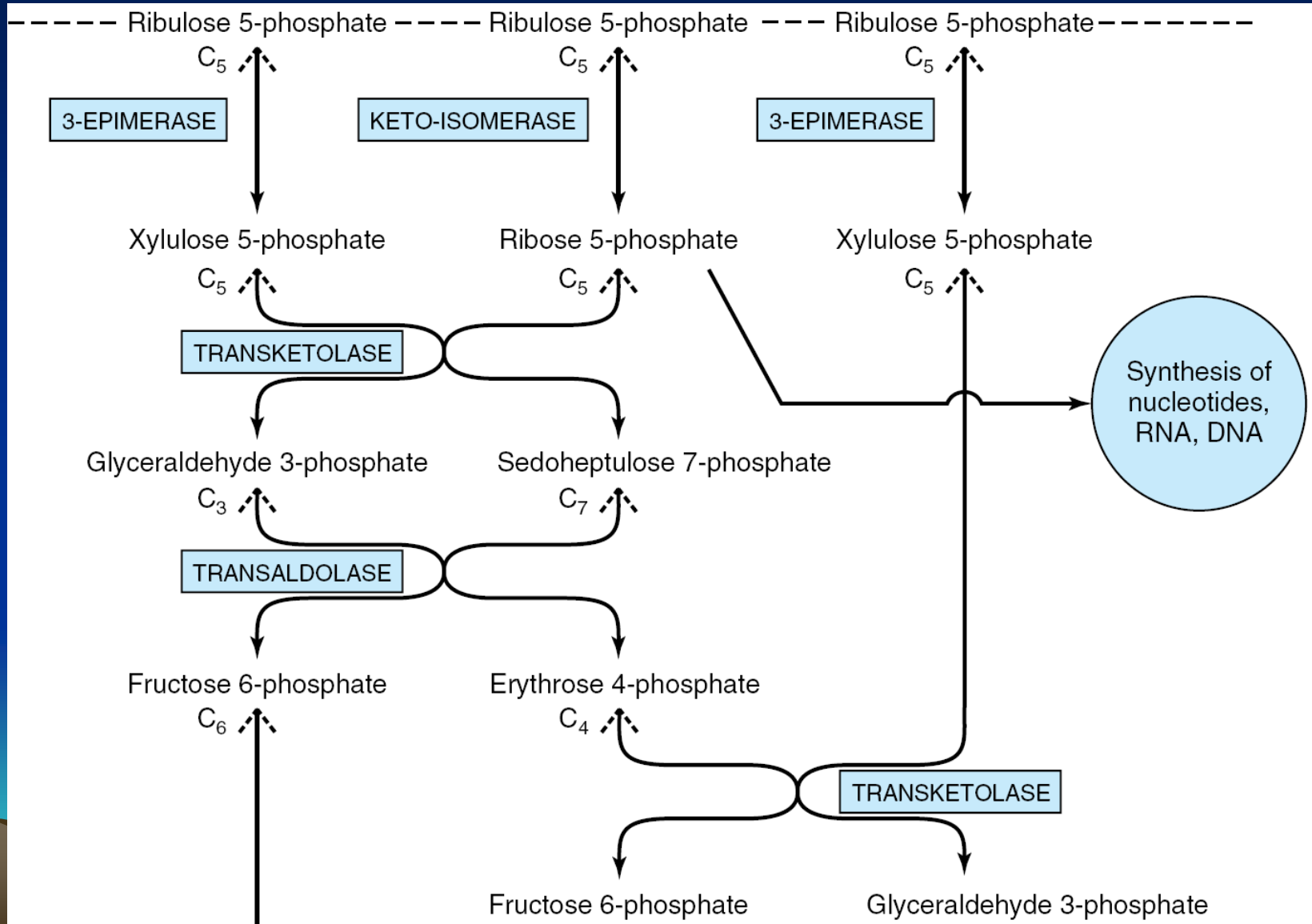
Pentose Phosphate Pathway

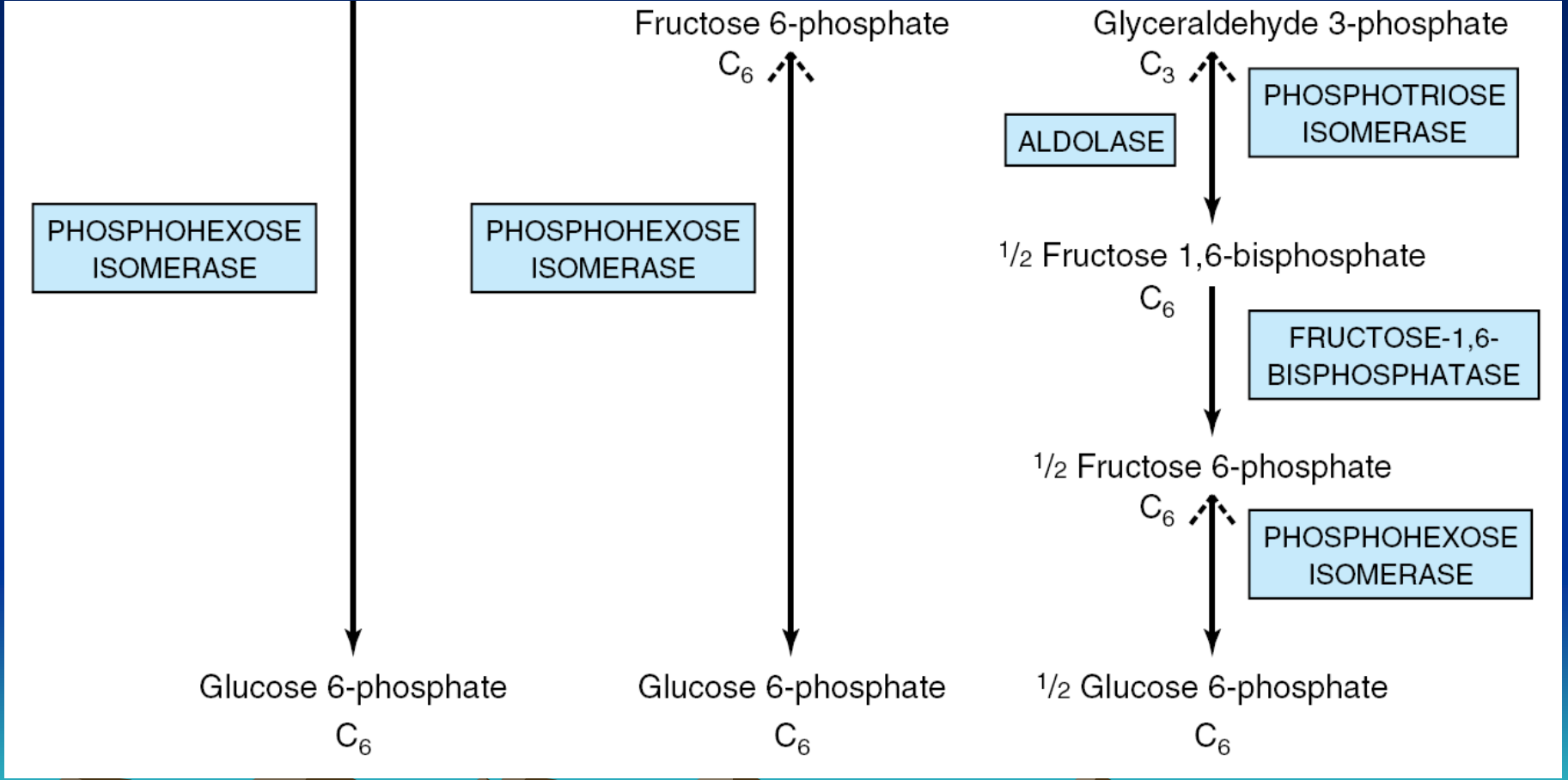


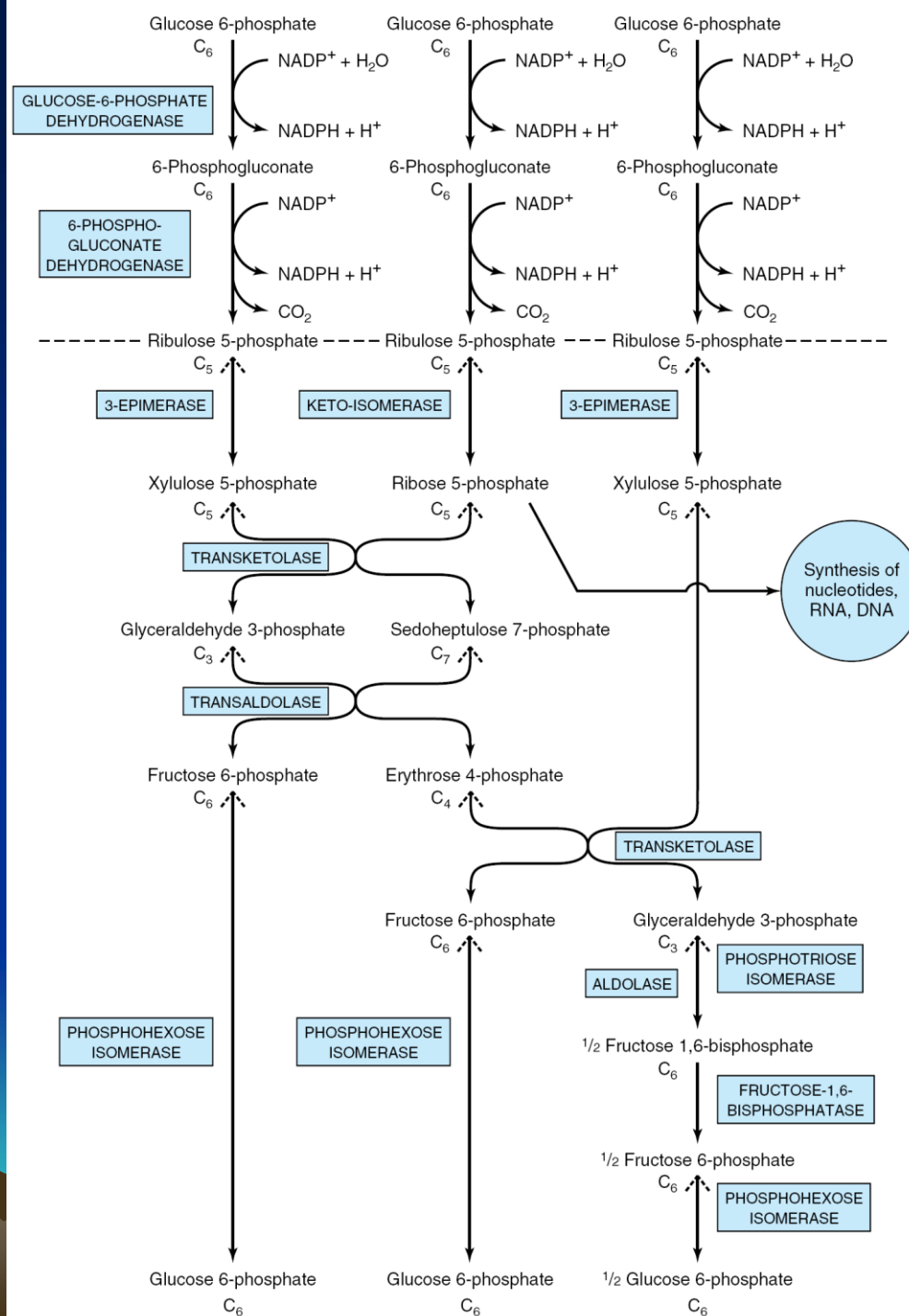
Oxidative phase

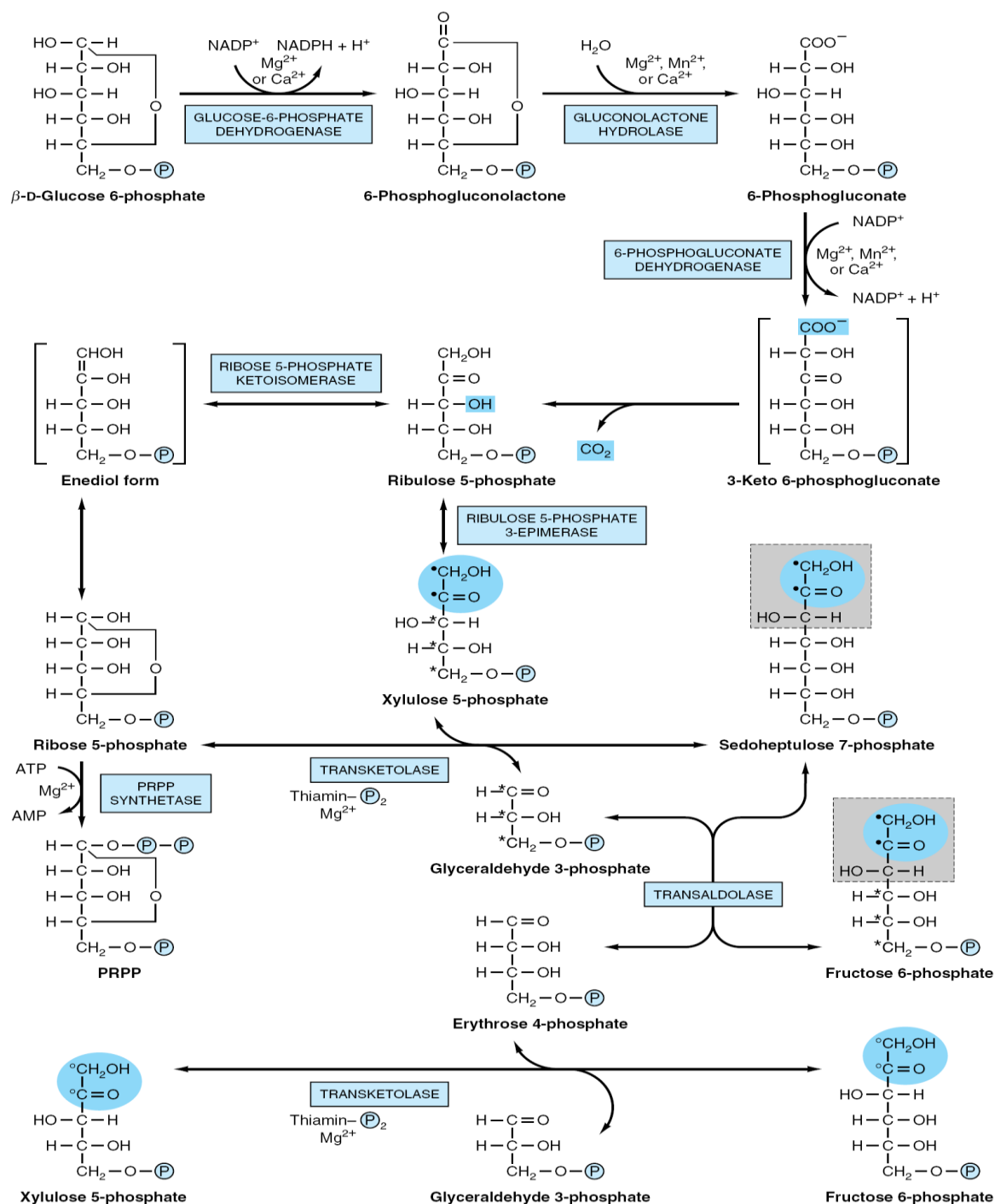


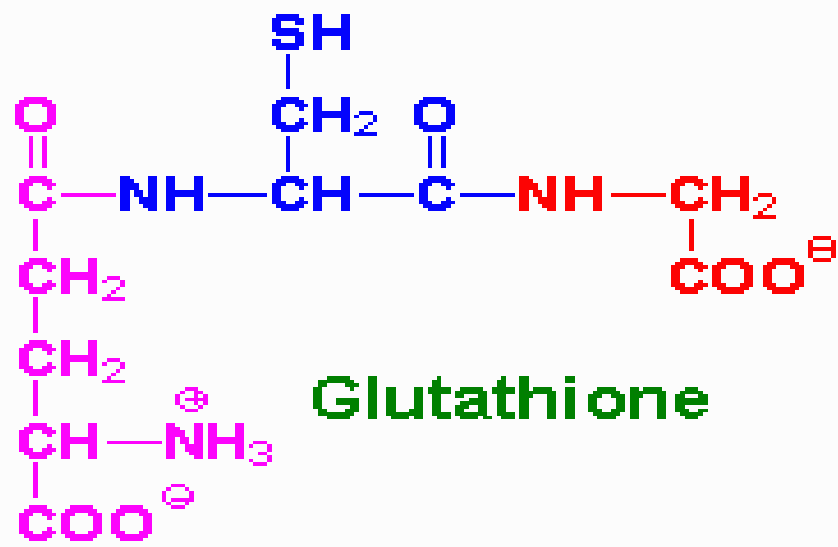
Non-Oxidative phase











Glutathione

Glutathione Role

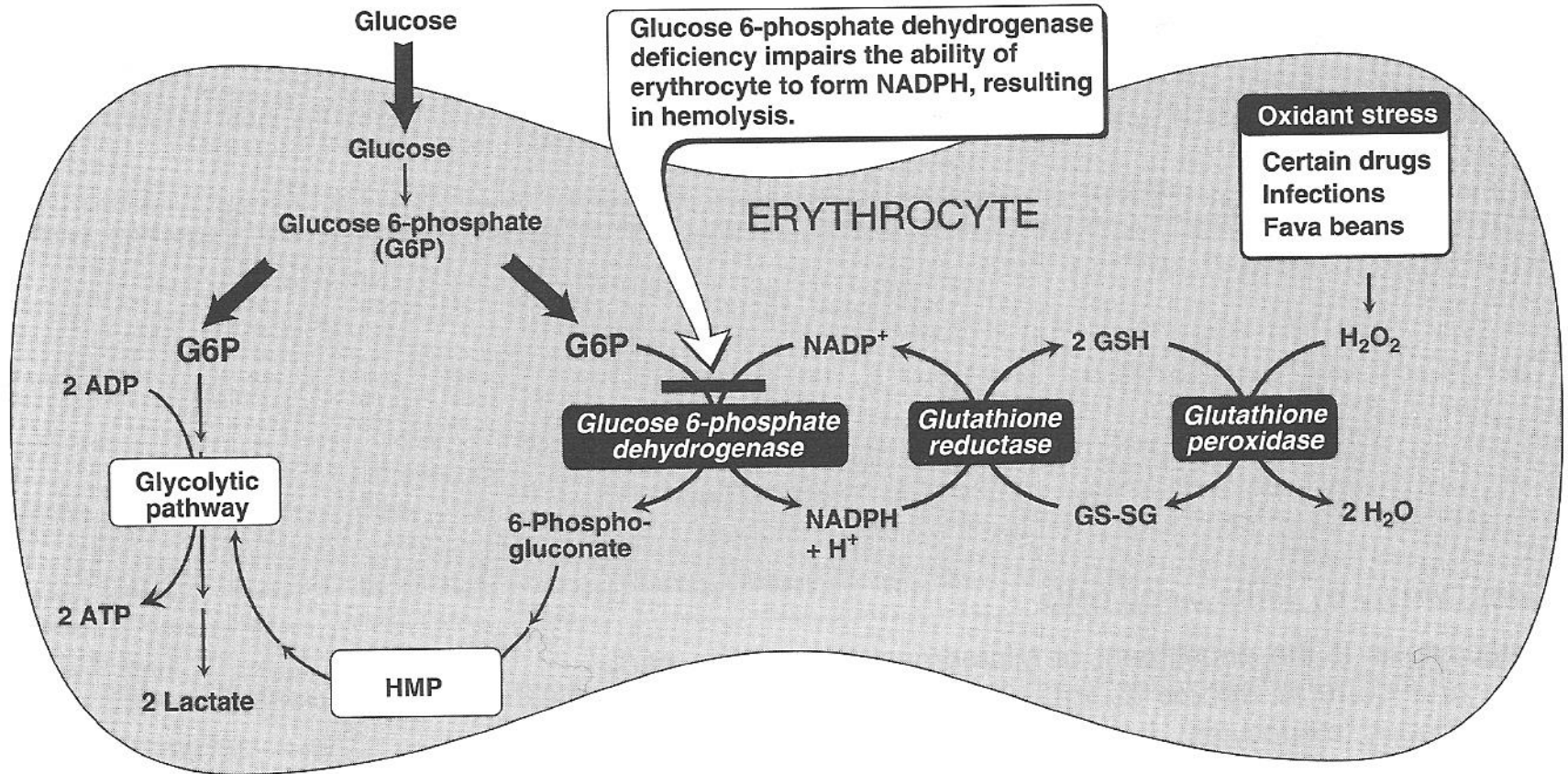


Figure 10.9

Pathways of glucose 6-phosphate metabolism in the erythrocyte.

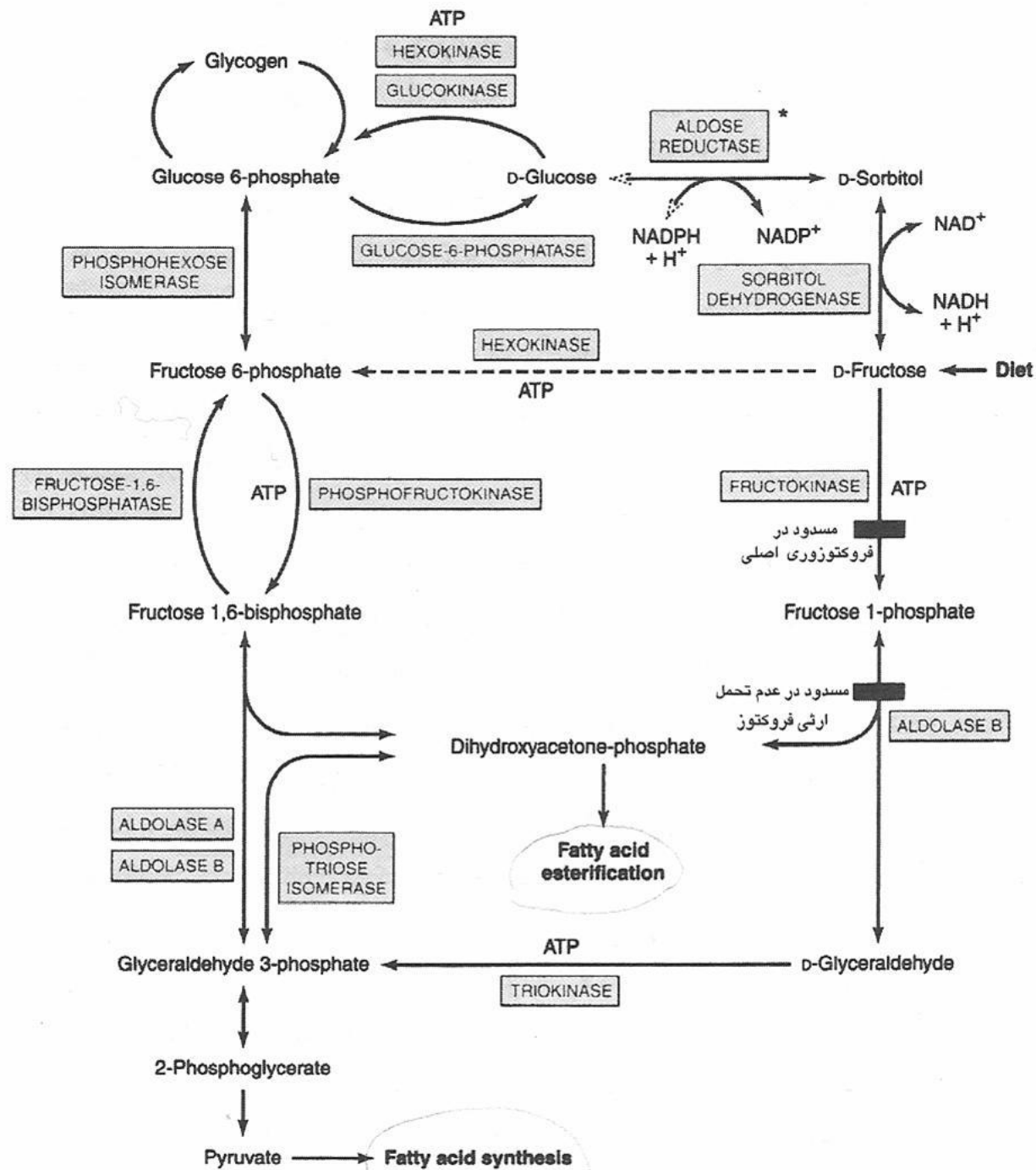
Enzymes of the pentose phosphate pathway and lipogenesis						
Glucose-6-phosphate dehydrogenase	↑	↓	Insulin			
6-Phosphogluconate dehydrogenase	↑	↓	Insulin			
"Malic enzyme"	↑	↓	Insulin			
ATP-citrate lyase	↑	↓	Insulin			
Acetyl-CoA carboxylase	↑	↓	Insulin?		Citrate, ¹ insulin	Long-chain acyl-CoA, cAMP, glucagon
Fatty acid synthase	↑	↓	Insulin?			

¹Allosteric.
²In adipose tissue but not in liver.



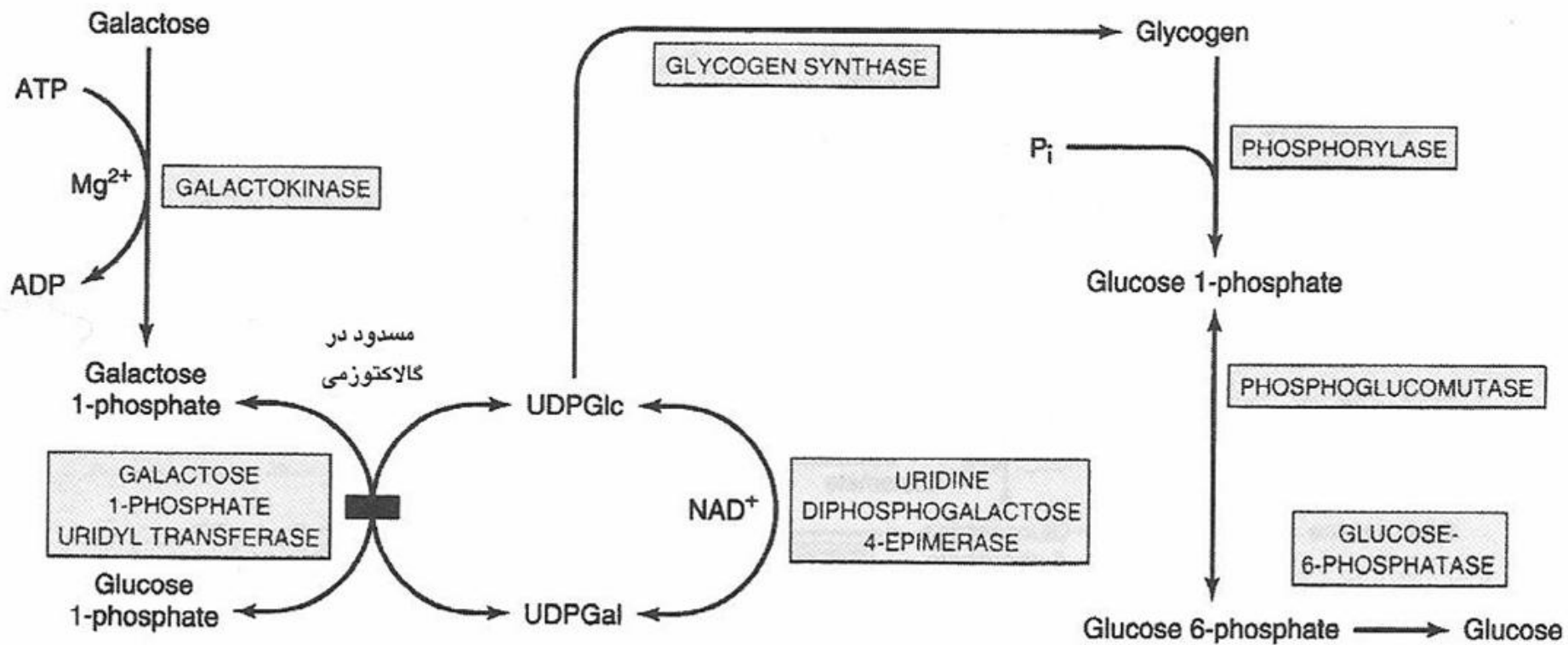
Fructose Metabolism





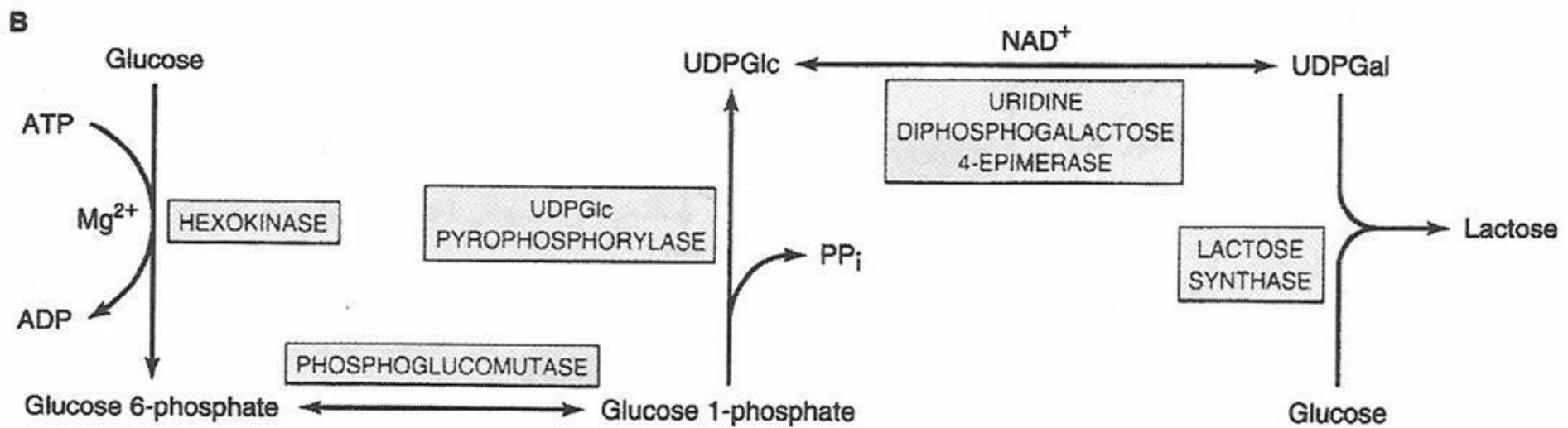
Galactose Metabolism





Lactose Metabolism





Cori Cycle

&

Glucose – Alanine Cycle



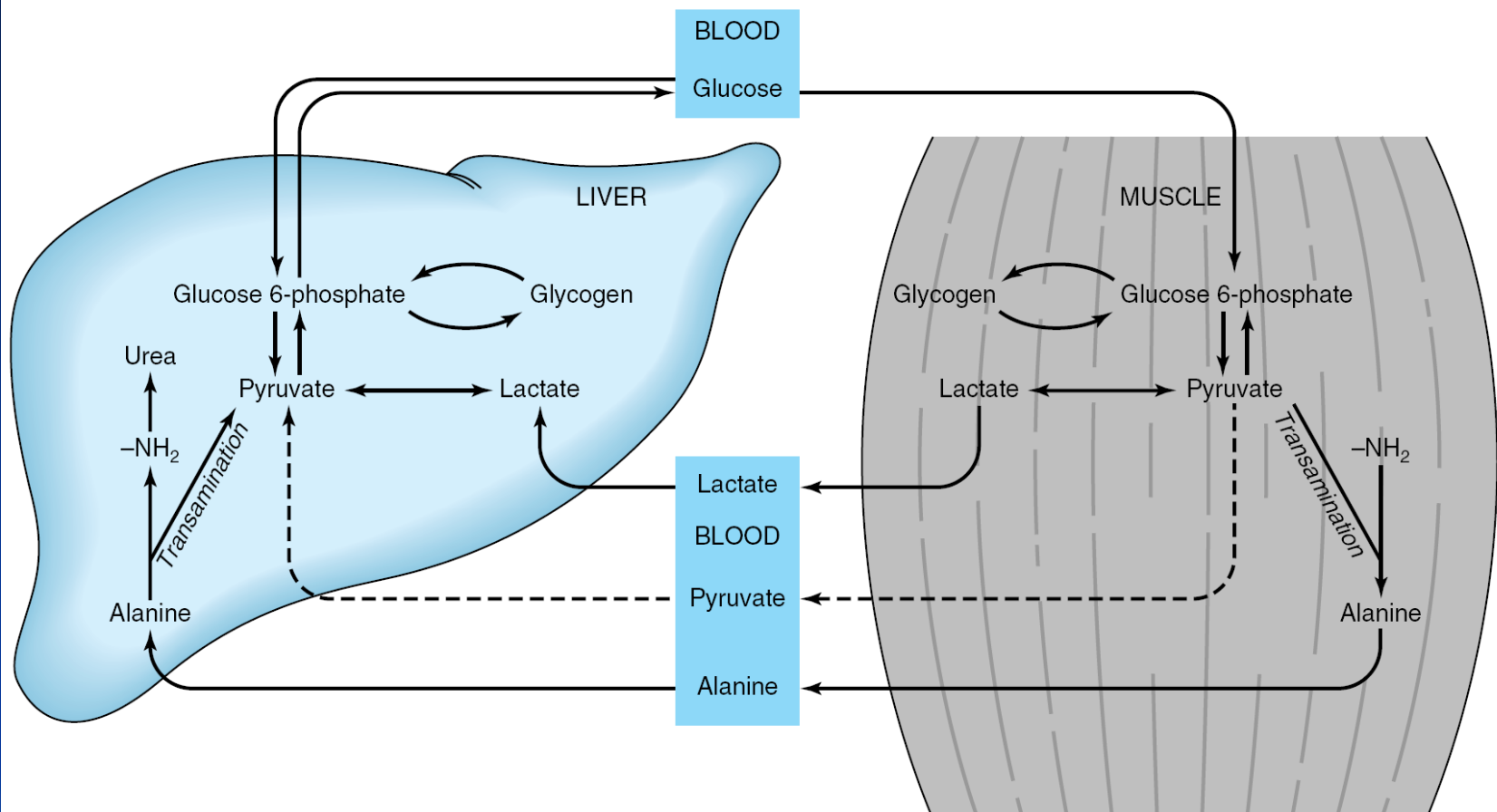


Figure 19-4. The lactic acid (Cori) cycle and glucose-alanine cycle.